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ADN35148 standard; protein; 229 AA. Human T1 receptor-like ligand II. US2003100048-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-AUG-1998.
(INCY-) INCYTE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-SEP-1999.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-SEP-2000.
(ALPH-) ALPHAGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                             29-MAY-2003.
(NIJJ/) NI J.
(GENT/) GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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Best Local Similarity
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RESULT 8
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protein - protein search, using sw model
n on: March 17, 2007, 11:54:03 ; Search time 217 Seconds
(without alignments)
516.393 Million cell updates/sec
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                                                                                                                                                                                                                                                                                               1 MGDKIWLPFPVLLLAALPPV......SAIQVYMLKSLFEDKRKSRT 229
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O1-JUL-2003.

(HUMA-) HUMAN GENOME SCI INC.

HUMA-) HUMAN GENOME SCI INC.

IOU.0%; Score 1177; DB 7; Length 229;

IET Match

O29;

O29;

O29;

O30;

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Pred. No. 9.6e-115;
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Angiogenesis protein AAA4 protein sequence (Fig 4).
WO200111086-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB87500 standard; protein; 229 AA.
Human II receptor (TIR)-like ligand II polypeptide.
US6586210-B1.
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26-FEB-1998.

(HUMA-) HUMAN GENOME SCI INC.

(ery Match 100.0%; Score 1177; DB 2;

(ery Match 100.0%; Pred. No. 9.6e-115;
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Pred. No. 9.6e-115;
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                                                                                                                                                                                                                                                                                                                                                                         Gapop 10.0 , Gapext 0.5
Searched: 2782304 segs, 489333398 residues
Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Listing first 1500 summaries
A Geneseq 200701:*
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Human T1-receptor-(T1R-)like ligand II.
US2002142461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW48335 standard; protein; 229 AA.
Human T1 receptor-like ligand II.
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Human T1R-like ligand II protein.
WO200142506-A1.
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geneseqp2003bs:*
geneseqp2004s:*
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geneseqp2002s:*
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MYCLOL-
(HUMA-) HUMAN GENOME SCI INC.
ery Match 100.0%;
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100.0%;
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Maximum DB seq length: 200000000
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1177
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Best Local Similarity
RESULT 2
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Best Local Similarity
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(RUBE/) RUBEN S M.
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                                                                                                                                                                                                                         Title:
Perfect score:
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Best Local S
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RESULT 1
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Run on:
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Length 229;
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ry Match
L Local Similarity 100.0%; Pred. No. 9.6e-115;
Length 229;
                                                                                                                                                                         Length 229;
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Human PRO733 (UNQ411) protein sequence SEQ ID NO:447.
WO200053756-A2.
(GETH ) GENENTECH INC.
100.0%; Score 1177; DB 7; 100.0%; Pred. No. 9.6e-115;
                                                                                                                    COCCT_2003.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 100.0%; Score 1177; DB 8;
                                                                                                                                                                                                                                                                                                                                               100.0%; Score 1177; DB 8; 100.0%; Pred. No. 9.6e-115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB34734 standard; protein; 229 AA.
Human secreted protein encoded by DNA clone vq20
WO200055375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99.5%; Score 1171; DB 2; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1171; DB 3;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW49100 standard; protein; 229 AA.
Human p24 vesicle trafficking protein Hp24-2.
WO9836068-A2.
                                                                                                                                                                                                                                      ADL91839 standard; protein; 229 AA. Human PRO83480 protein SEQ ID NO:60.WO2004024076-A2.
                                                                                                                                                                                                                                                                                                                                                                                                             ADR03638 standard; protein; 229 AA. Human TIR-like ligand II protein. US2004146501-A1.
                                                                ADI34089 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   229 AA
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                                                                                  Human T1R-like ligand II protein.
US2003198618-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY41754 standard; protein; 22
Human PRO733 protein sequence.
WO9946281-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 99.6%;
RESULT 12
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Best Local Similarity 99.6%;
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Local Similarity 99.6%;
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99.5%; Score 1171; DB 3; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                        99.5%; Score 1171; DB 6; Length 229; 99.6%; Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.5%; Score 1171; DB 6; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                Length 229;
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                                                                                                                                                                           AB017820 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US20030312156-Ai.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU72262 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2002192706-AI.
                                                                                                                                                                                                                                                                                                          ABO25256 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003050239-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABU84942 standard; protein; 229 AA.
Human secreted and transmembrane PRO polypeptide #18.
US2002177553-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-NOV-2002.
(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 6;
                                                                                                                                                                                                                                                           Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                Score 1171; DB 4;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABUS9855 standard; protein; 229 AA.
Novel secreted and transmembrane protein PRO733.
US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #205.
US200336180-A1.
                                              AAU12376 standard; protein; 229 AA. Human PRO733 polypeptide sequence. WO200140466-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                     ABUB1074 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003004311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABU61140 standard; protein; 229 AA
                                                                                                                                                                                                                                                         99.5%;
                                                                                                                              99.5%;
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US2002169284-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                 (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 23
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Query Match
Best Local Similarity
                                                                                                                                           Local Similarity
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                               RESULT 15
                                                                                                                                                               RESULT 16
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Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003022328-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABUG7050 standard; protein; 229 AA.
Human secreted/transmembrane, PRO, protein SEQ ID
US2003032155-A1.
                                                                                                      20-FEB-zuus.
(GETH ) GENENTECH INC.
pry Match 99.5%; Score 1171; DB 6;
rear eimilarity 99.6%; Pred. No. 4.1e-114;
Best Local Similarity 99.6%; Pred. No. 4.1e-114; RESULT 24
                                                                                                                                                                                                                                                                                      Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                         ABO25045 standard; protein; 229 AA.
Human secreted/transmembrane protein (PRO) #205.
US2003036179-A1.
                                                                                                                                                                                                                                                                                                                                                           Human p24 vesicle trafficking protein Hp24-2. US2002132999-A1.
                                                                                                                                                                                          ABUB0409 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733.
US2003004102-A1.
                                                                                                                                                                                                                                                                                                                                           ABG72289 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA76360 standard; protein; 229 AA Human PRO polypeptide #205.
US2003073212-A1.
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0S2003054517-A1.
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02-0AN-2003.
(GETH ) GENENTECH INC.
99.5$;
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ry Match 99.5%;
t Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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US2003049816-A1.
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OJ-WAY-2-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
                                                                                                                                                                       GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
(ery Match 99.5%; Score 1171; DB 6; Length 229;
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                                                                                                                                                Novel human secreted and transmembrane protein PRO733 US2003082711-A1.
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Novel human secreted and transmembrane protein PRO733
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                               Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
            ADB27959 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082704-A1.
                                                                                                                                                                                                                                                  ADB16002 standard; protein; 229 AA. Human PRO polypeptide #205. US2003087350-A1. GB-MAY-2003. (GETH ) GENENTECH INC.
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US2003082705-Al.
                                                                                                                                                                                                                                                                                                                                                                       ADA47788 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003073215-Al.
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US2003082763-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 229 AA
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                                                                                                                                ADA86438 standard; protein; 229 AA
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01-MAY-2003.
(GETH ) GENENTECH INC.
99.5%; SC
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US2003068795-A1.
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US2003068794-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 37
ID ADAG7583 standard; pr
DE Human PRO polypeptide
PN US2003068795-A1.
PD 10-APR-2003
PA (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 39
                                                                                               Best Local Similarity RESULT 34
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Best Local Similarity
RESULT 42
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RESULT 33
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01-MAY-2003.
(GETH) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
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Pred. No. 4.1e-114;
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ADA87541 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003087345-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003050241-A1.
                                                                                                                                                                                                                                                                             ADA91835 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082694-A1.
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Novel human secreted and transmembrane protein PRO733
US2003082691-A1.
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                        Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                              Human PRO polypeptide #205.
US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                         ADB14898 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087351-A1.
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17.APR-2003.
(GETH ) GENENTECH INC.
99.5%; S
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(GETH ) GENENTECH INC.
99.5%;
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PA (GETH ) GENENTECH INC.

QUETY MATCh 99.5%;

Best Local Similarity 99.6%;
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1 (GETH) GENENTECH INC.

Query Match

Best Local Similarity 99.6%;
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99.6%;
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(GETH ) GENENTECH INC.
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RESULT 51
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Best Local Similarity
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                                                                                                                           RESULT 43
                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 45
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17-APR-2003.
(GETH ) GENENTECH INC.
US2003073214-A1.
                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC. (GETH ) GENENTECH INC. 99.5%; Score 1171; DB 6; Length 229; ery Match 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OL-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
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Pred. No. 4.1e-114;
                                                                                                                                                                                Score 1171; DB 6; Length 229;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                           Length 229
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Novel human secreted and transmembrane protein PRO733.
US2003082708-A1.
Novel human secreted and transmembrane protein PRO733. US2003044945-A1.
                                                                                                     ABO19711 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003050240-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA85334 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733. US2003082695-A1.
                                                         Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99.5%; Score 1171; DB 6; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                      Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                           ADA12647 standard; protein; 229 AA.
Human secreted/transmembrane polypeptide PRO733.
US2003055216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB24769 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077713-A1.
                                                                                                                                                                                                                                                                                                                                                ADA74536 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003068798-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA82293 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003082701-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA75256 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003073216-A1.
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Human PRO polypeptide #205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lery Match GENENTECH INC.
Best Local Similarity 99.6%; Pre RESULT 56
ID ADA82293 standard; prot PN US200308270**
PD US200308270**
PD 01-MP**
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17-APR-2003.
(GETH ) GENENTECH INC.
99.5%; S'
                                                         99.5%;
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Best Local Similarity 99.6%;
RESULT 54
                                                                                                                                                 13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                        20-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                      28-A1.
2003.
2003.
2003.
2004.
2007.
Best Local Similarity SRESULT 55
ID ADB24769 stand*
DB Human PRO r
PN US2003^c
PD 24.
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2003.

Lift ) GENENTECH 1

STY MATCH

Best Local Similarity PRESULT 57

ID ADA72256 stand*

DE Human PRO T

PN US20037*

PD 17
      (GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 60
ID ADB30038
                                                                                                                                                                                           Best Local Similarity RESULT 53
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IS-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
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Query Match ,99.5%; Score 1171; DB 6; Length 229; Best Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                           Length 229;
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(GETH) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
                                                                                                                                                                                                                                                  Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 229;
                                    ID ADA80566 standard; protein; 229 AA.

DE Human PRO polypeptide #205.

PN US2003082761-A1.

PD 01-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 62.
                                                                                                                                                                                                                                                    Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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US2003077714-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                      ADB25329 standard; protein; 229 AA. Human PRO polypeptide SEQ ID NO 410. US2003077715-A1.
                                                                                                                                                                                                                                                                                    Human PRO polypeptide #205.
17-App.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA93505 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003077721-A1.
                                                                                                                                                                      ADA75808 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082703-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 229 AA
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17-APR-2003.
(GETH ) GENENTECH INC.
Watch '13-11LY 99.5%;
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24-APC-2003.
(GETH ) GENENTECH INC.
MATCh ''1arity 99.6%;
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ADB26855 standard; protein;
Human PRO polypeptide #205.
US2003092147-Al.
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US2003096386-A1.
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(GETH ) GENENTECH INC.
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.-2003.
.-aery Match
Best Local Similarity
RESULT 68
ID ADA61070 stand*
DE Homo sapie*
PN US2003*
PP 13
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Best Local Similarity
RESULT 64
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Best Local Similarity
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Best Local Similarity
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Best Loc
RESULT 86
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24-APR-2003.
(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 6; Length 229;
Lery Match 99.6%; Pred. No. 4.1e-114;
                                                                                                                                     Score 1171; DB 6; Length 229;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane polypeptide #70.
US2003049633-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA86990 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082709-A1.
01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB21788 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003082765-A1.
                                                                                                                                                                                                                                                                                                                                                 Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                  ADA96546 standard; protein; 229 AA. Human PRO polypeptide #205. US2003002690-A1. Ols-MAY-2003 (GETH ) GENENTECH INC.
                                                                                                                                                                   Human PRO polypeptide #205.
US2003082702-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #205.
US2003068797-Al.
                                                                                                                                                                                                                                                                               ADA95994 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082759-A1.
                                                                                                                                                                                                                                                                                                                                                                                    ADB26303 standard; protein; 229 AA. Human PRO polypeptide #205. US2003082760-A1.
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US2003077710-A1.
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01-MAY-2003.
(GETH ) GENENTECH INC.
***rch 99.5%; Sc
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01-MAY-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; S/
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01-MAY-2003.
(GETH ) GENENTECH INC.
Watch 99.5%; S/
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01-MAY-2003.
(GETH ) GENENTECH INC.
99.5%;
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(GETH ) GENENTECH INC.
99.5%;
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
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                                                                                                                                                                                                                                                                                                                                         query Match
Best Local Similarity
RESULT 73
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Best Local Similarity
RESULT 74
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Best Local Similarity
RESULT 76
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                                 Query Match
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Length 229;
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        Length 229;
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                                                 ADA88093 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082700-A1.
                                                                                                                                                                                ADA46481 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003054516-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       אראפינה standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003073213-A1.
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      99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                  99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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US2003022239-Al.
310-DAN-2003.
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US2003059909-A1.
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                                                                                                                                                                                                                                                                                                            ADB28511 standard, protein, 229 AA Human PRO polypeptide #205. US2003082699-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB29063 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082706-A1.
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99.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                20-MAR-2003.
(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
Best Local Similarity
RESULT 79
                                                                                                                                              Best Local Similarity RESULT 80
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Best Local Similarity
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RESULT 88
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                                                                                                                                    Query Match
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Human PRO polypeptide #205.
US2003082762-Al.
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                                                                              Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                          ADA92387 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082712-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB38702 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB30150 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733
US2003087347-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB66622 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082689-A1.
                                                                                                                                                                                                            99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                  Human PRO polypeptide SEQ ID NO 410.
US2003077712-A1.
ADA67031 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003068793-A1.
                                                                                                                          ABB22892 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003077711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB15450 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003087352-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99.5%;
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                               PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB89702 standard; protein;
Human PRO polypeptide #205.
US2003082698-A1.
                                                                                                                                                      2003.

2003.

2003.

2ry Match

Beet Local Similarity 5
RESULT 90
ID ADB23665 stand**
DE Human PRO r.
PN US2003 r.
PD 24-*
                                                                                                                                                                              24-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                           24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
RESULT 91
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Best Local Similarity
RESULT 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-MAY-2003
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Length 229;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                 ADB41158 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB77370 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082696.A1.
01-MAY-2003.
                                                                                                                        Novel human secreted and transmembrane protein PRO733 US2003082764-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human PRO polypeptide SEQ ID NO 410.
US2003077717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PRO polypeptide SEQ ID NO 410.
US200307719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB33975 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410
                                                                                                        protein; 229 AA.
                                                                                                                                                                                                                                                protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB86765 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082697-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 229 AA
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-2003.
) GENENTECH INC.
99.5%; Sr
JS2003002.
01-MXY-2003.
(GETH ) GENENTECH INC.
99.5%;
                                                                                                                                                                                                                                                                                               06-MAR-2003.
(GETH ) GENENTECH INC.
Query March 99.5%;
Best Local Similarity 99.6%;
                                                                                                                                                        01-MAY-2003.
(GETH ) GENENTECH INC.
2ry Match 99.5%;
st Local Similarity 99.6%;
                                                                                                                                                                                                                                             ADB73953 standard; proteir
Human PRO polypeptide #70.
US2003045462-A1.
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US2003083248-A1.
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(GETH ) GENENTECH INC.
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                                                                Best Local Similarity RESULT 98
                                                                                                        ADB39535 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB76669 standard;
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Query Match
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                                                             RESULT 116
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                                            99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                             Score 1171; DB 7; Length 229;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                Score 1171; DB 7; Length 229;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                       ADB46578 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082692-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-mar. 2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 7;
ery Match 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                               Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC44095 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003054986-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC61819 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003054405-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC66919 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003060406-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC61855 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003049684-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC69043 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003064407-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC63103 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003068648-A1.
                                                                                                                                                                                                    ADB36183 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077720-A1.
24-APR-2003.
(GETH ) GENENTECH INC.
24-APR-ACC.
39.5%; Score 117
best Local Similarity 99.6%; Pred. No.
                                                                             Human PRO polypeptide SEQ ID NO 410.
US200307718-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lery Match 10 GENENTECH INC.
Best Local Similarity 99.6%; Prec RESULT 114
ID ADC69043 standard; prof DE Human secreted/from PD 03.APP
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20-MAR-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; SC
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13-MAR-2003.
(GETH ) GENENTECH INC.
Watch 99.5%; Sc
                                                                                                                                                           99.5%;
US2003077716-A1.
24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                24-APR-2003.
(GETH ) GENENTECH INC.
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Sert Local Similarity
RESULT 115
ID ADC63103 Str
DE Humar
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Best Local Similarity
REULT 110
ID ADC44095 standard; pr
DE Human secreted/transn
PN US2003054986-A1.
PD 20-WAR-2003.
PA (GETH ) GENENTECH INC
                                                     Best Local Similarity RESULT 107
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Best Local Similarity
RESULT 112
                                                                                                                                                                       Best Local Similarity
RESULT 108
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RESULT 109
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Best Local Similarity
RESULT 111
                                                                                                                                                                Query Match
                                              Query Match
                                                                                                                                                                                                                                                                                Query Match
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Length 229;
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Novel human secreted and transmembrane protein Seg ID410.
US2003087365-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC71998 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092107-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human secreted and transmembrane protein PRO733. USAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human secreted and transmembrane protein PRO733 US2003092106-A1.
                               Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                            Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                      Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC42112 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003104998-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                       ADC41488 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003072745-A1.
                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, PRO733.
US2003073131-Al.
                                                                          ADC68168 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003069178-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, PRO733. US2003073624-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 229 AA.
                                                                                                                                                                                                                                                         US20030,17-ARR-2003.
(GETH ) GENENTECH INC.
99.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USZUCZZ
15-WAY-2003.
(GETH ) GENENTECH INC.
WALCh '-'1arity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 99.6%;
RESULT 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Ouery March 99.5%;
Best Local Similarity 99.6%;
10-APR-2003.

A (GETH) GENENTECH INC.

Query Match 99.5%;

Best Local Similarity 99.6%;
                                                                                                                             PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 99.5%;
Best Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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PA (GETH ) GENENTECH INC.
Query Match 99.5%;
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 123
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC62479 standard;
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Length 229;

Length 229;

Length 229;

Length 229;

us-10-692-730-2.spdi.rag

RESULT 126

RESULT

RESULT 125

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Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003087354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC80595 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092103-A1.
                                                 ADD01235 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003092104-A1.
                                                                                                                                                                                           ADC90227 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087348-A1.
Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                         Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.
99.5%; Score 1171;
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US200199471-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #205.
1052003194776-Al.
                                                                                                                                                                                                                                                                                                                                     ADC69646 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194770-Al.
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US2003194773-A1.
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Human PRO polypeptide #205.
US2003194774-A1.
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08-MRY-2003.
(GETH ) GENENTECH INC.
MAtch '1-rity 99.6%;
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15-MAY-2003.
(GETH ) GENENTECH INC.
MATCh ''ATTICY 99.5$;
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16-OCT-2003.
(GETH ) GENENTECH INC.
MATCh --1arity 99.6%;
99.5%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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ry Match 99.5%;
c Local Similarity 99.6%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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              Best Local Similarity RESULT 134
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Best Local Similarity
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 Query Match
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Pred. No. 4.1e-114;
                  Length 229;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human secreted and transmembrane protein Seg ID410. US200387363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC53590 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC59113 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC55991 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087360-A1.
                                                                  ADC57338 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADCS8561 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410
US2003087346-A1.
                                                                                                                                                                                                            ADC60529 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087367-A1.
                                                                                                                                                                                                                                                                                                                                                      ADC51004 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087361-A1.
                  Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human PRO polypeptide #205.
US2003087362-A1.
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08-MAY-2003.
(GETH ) GENENTECH INC. 99.5%;
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                99.5%;
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RESULT 128
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(GETH) GENENTECH INC.
PRY MATCH 99.5%;
it Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 131
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Best Logal Similarity
RESULT 132
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Best Local Similarity
RESULT 130
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                Query Match
Best Local Similarity
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RESULT 129

Length 229;

Length 229;

Length 229;

Length 229;

Length 229;

DB 7;

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vuery Match
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 150
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                               ADC80043 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087358-A1.
                                                                                                                                                                                                                                                                                                   ADD41225 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aurosisse standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US203203437-A1.
99.6%; Pred. No. 4.1e-114;
                                                                                                                 Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                  ADD09512 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194775-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD52364 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD53104 standard; protein; 229 AA. Human PRO polypeptide #205. US2003194792-A1.
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                                                               US200500...
08-MAY-2003.
(GETH ) GENENTECH INC.
99.5%;
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16-007-2003.
(GETH ) GENENTECH INC.
99.5%;
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30-OCT-2003.
(GETH ) GENENTECH INC.
99.5%;
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US2003203430-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                                                                                Query Match
Best Local Similarity
RESULT 144
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                Length 229;
Novel human secreted and transmembrane protein PRO733. US2003203432-A1. 30-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB32351 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003194765-A1.
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Pred. No. 4.1e-114;
                                                                                               Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                            Best Local Similarity 99.6%; Fred. No. 4.1e-1
RESULT 153
ID ADE49481 standard; protein; 229 AA.
DE Human secreted/transmembrane protein, PRO733.
PN US2003096744-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, PRO733. US200334-A1.
                                                                                                                                                                                                                                                                     ADD92544 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199030-A1.
23-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE04054 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199057-A1.
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Human PRO polypeptide #205.
US2003199055-A1.
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Human PRO polypeptide #205.
US2003199056-A1.
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US2003203428-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-2003.
(GETH ) GENENTECH INC.
Query Match
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                                                                                                                                                                                                                                 Best Local Similarity RESULT 154
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Best Local Similarity
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RESULT 161
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ID ADE35535 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-2003
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                                                                                                Query Match
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   RESULT 152
ID ADD542
DE Novel
PN US2003
PD 30-OCT
PA (GETH
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Human secreted/transmembrane protein, PRO733. US2003194781-A1.

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Beet Local Similarity 99.5%; Score 1171; DB 7; Length 229;
Beet Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 164
ID ADE17860 standard; protein; 229 AA.
DE Human PRO polypeptide #205.
PN US2003199023-A1.
PA //--
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                                                                                         Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE33455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
10/22003194/67-A1.
16-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE34007 standard, protein, 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194791-A1.
                                                                                        Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                   Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
ADE16649 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203435-A1.
                                                                                                                                           ADD73264 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD91992 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003199053-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD80059 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207417-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD93096 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194768-A1.
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                                  US2003201.
30-OCT-2003.
(GETH ) GENENTECH INC.
99.5%;
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                                                                                                                                                                                           PD 30-OCT-2003.

PA (GETH ) GENENTECH INC.

Query Match 99.5%;

Best Local Similarity 99.6%;

RESULT 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-A1.
2003.
21.H J GENENTECH 1.
2ry Match
Best Local Similarity 3.
RESULT 165
ID ADD91992 standal
DE Human PRO ri
PN US20031
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(GETH ) GENENTECH INC
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                                                                                                   Best Local Similarity RESULT 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
RESULT 166
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Best Local Similarity
RESULT 170
ID ADD72622 standard;
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Best Local Similarity
RESULT 169
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(GETH ) GENENTECH INC.
iry Match
99.5$; Score 1171; DB 7; Length 229;
rry Match
oral Similarity 99.6$; Pred. No. 4.1e-114;
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                                                       Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003194766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                     Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
                                                                                                                                                                                                                                                                                                                                                                                                 ADE43160 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199033-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD95949 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199059-A1.
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Human PRO polypeptide #205.
US2003199064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD78953 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003203429-Al.
                                                                                                          ADE19516 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199025-A1.
                                                                                                                                                                                                                                                         protein; 229 AA
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US2003202.
30-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S'
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23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S.
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(GETH) GENENTECH INC.
Query Match 99.5%;
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
PRY MATCh 99.5%;
st Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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(GETH) GENENTECH INC.
10-GETY MATCh 99 5%;
Best Local Similarity 99.6%;
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US2003199026-A1.
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US2003199032-A1.
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(GETH ) GENENTECH INC.
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                                                                      Best Local Similarity RESULT 171
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GENENTECH INC.
 06-NOV-2003
(GETH ) GEN
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                                                               RESULT 189
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         30-OCT-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 7; Length 229;
                                                                                                                                                                    Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207384-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG21560 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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RESULT 185
ID ADF47288 standard; protein; 229 AA.
DE Human Secreted/transmembrane protein, PRO733.
PN US2003195333-A1.
                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #205.
US2003199031-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #205.
US2003194777-Al.
                                                                                            protein; 229 AA.
                                                                                                                                                                                                                ADD89639 standard; protein; 229 AA. Human PRO polypeptide #205. US2003199028-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE04722 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE95520 standard; protein; 229 AA.
Human NOVX4c protein.
WO2003050245-A2.
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23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S.
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23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S'
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16-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S.
                                                                                                                     US200sco...
06-NOV-2003.
(GETH ) GENENTECH INC.
99.5%;
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23-OCT-2003.
(GETH ) GENENTECH INC.
"" OF Ch. "" OF S$;
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                                                                                                         Human PRO polypeptide #205.
US2003207418-A1.
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US2003199034-A1.
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(GETH ) GENENTECH INC.
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(CURA-) CURAGEN CORP
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                                                      Best Local Similarity
RESULT 180
                                                                                                                                                                    Ouery Match
Best Local Similarity
RESULT 181
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 182
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Best Local Similarity
RESULT 183
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RESULT 184
ID ADE92851 standard;
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                                                                                          ADD80611 standard;
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.US2003203433-A1
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                                            Query Match
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Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
   Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207381-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH55892 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207379-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI64111 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
052003207385-A1.
06-NOV-2003.
99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
                                                                                                                             99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                        Human secreted/transmembrane protein, PRO733 US2003216561-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI61124 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003077700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG80048 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207372-A1.
                                                 229 AA
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                                         Human PRO polypeptide #205.
082003207370-A1.
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                                                                                                                                                                                                                                                                                                         ADG53044 standard; protein;
                                                                                          06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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 Query Match
Best Local Similarity
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Best Local Similarity
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RESULT 190
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Best Local Similarity
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RESULT 192
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Length 229;

Length 229;

us-10-692-730-2.spdi.rag

RESULT 199

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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                      ADC81147 standard, protein, 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003092115-A1.
99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                               Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003104536-A1.
05-JUN-2003.
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(GETH.) GENENTECH INC.
99.5*; Score 1171;
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US2003092110-A1.
                                                                                                                                                                                                                                                                    ADD76595 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003100087-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADD87959 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003092113-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE75811 standard; protein; 229 AA Human PRO polypeptide #205.
US2003211571.A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE23387 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003092108-A1.
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15-MAZ-2003.
(GETH ) GENENTECH INC.
99.5%;
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(GETH ) GENENTECH INC.
1-V Match 99.5%;
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Local Similarity 99.6%;
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Query Match 99.5%;
Best Local Similarity 99.6%;
RESULT 213
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1 (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 99.6%;
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Human PRO polypeptide #205.
US200320340-A1.
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1 (GETH) GENENTECH INC.

10-OCTY MATCh

10-OCTY MATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 99.5%;
Best Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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US2003092111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAY-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 215
               Best Local Similarity RESULT 207
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     Query Match
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                      Length 229
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003087357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM82590 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003087355-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003087385-A1.
                                                                                                                                                                                                                                                                                        ADI63559 standard; protein; 229 AA.
Novel human secreted and transmembrane protein FRO733.
US2001207387-A1.
(GETH ) GENENTECH INC.
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Novel human secreted and transmembrane protein PRO733
US2003207377-A1.
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Novel human secreted and transmembrane protein PRO733
US2003087353-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003087356-A1.
                                                                                             ADI65060 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003207386-A1.
                                                                                                                                                                                                                      Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                           Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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08-MAY-2003.

(GETH ) GENENTECH INC.

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08-MAY-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; SC
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08-MAY-2003.
(GETH ) GENENTECH INC.
99.5%;
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LH ) GENENTECH IN
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Best Local Similarity 9
RESULT 206
ID ADNI4885 standa-
DE Novel humar
PN US2003^C
PD 08
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 200

ID ADH81973 stands

DE Novel humar

PN US20032

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(GETH ) GENENTECH INC.
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RESULT 202

ID ADM82590 stander

DE Novel humar

PN US2003r

PD 08-7
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(GETH ) GENENTECH INC.
  (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 203
                      Query Match
Best Local Similarity
RESULT 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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RESULT 201 ID ADH814 DE NOVEL PN US200: PD 06-NO' PA (GETH

Length 229;

Length 229;

DB 8;

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Length 229;
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    Length 229;
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                                                                                                                             Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
    99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                          ADF46010 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195148-A1.
                                                ADF40214 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003198994-Al.
23-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aurana secreted/transmembrane protein, PRO733.012004055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein, PRO733. US2003199021-A1.
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102203203402-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GETH ) GENENTECH INC. 99.5%; Score 1171;
                                                                                                                                                                                                                                                                                                    protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE95293 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199052-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 229 AA
                                                                              US200312C1.
23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S'
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23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%;
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Local Similarity 99.6%;
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Local Similarity 99.6%;
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Local Similarity 99.6%;
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(GETH) GENENTECH INC.
PY MATCH 99.5%;
it Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE93403 standard; protein;
Human PRO polypeptide #205.
US2003199060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 99.5%;
Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                   ADE94741 standard; protein;
Human PRO polypeptide #205.
US2003199027-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                             ADE91152 standard; protein;
Human PRO polypeptide #205.
US2003199061-A1.
                                                                                                                                                                                                                         16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                           Best Local Similarity
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    Query Match
Best Local Similarity
RESULT 222
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ID ADF24406 standard;
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                                                                                                              Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003130181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF61522 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195345-A1.
                                                                                                                                                       ADE89273 standard; protein; 229 AA. Human PRO polypeptide #205. US2003199062-A1. CGETH ) GENENTECH INC.
                           ADD87407 standard; protein; 229 AA. Human PRO polypeptide #205. US2003203439-A1. 30-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                 ADE18412 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                           ADE88721 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199054-A1.
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(GETH ) GENENTECH INC.
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BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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SHELTON D L.
STEWART T A.
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WILLIAMS P M.
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KLJAVIN I J.
KUO S S.
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RESULT 219
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PAONI N F.
Best Local Similarity
RESULT 216
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FERRARA N.
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Best Local
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Query Match

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Length 229;
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Pred. No. 4.1e-114;
                                                                                                            Length 229;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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US2001199435-A1.
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Human secreted/transmembrane protein, PRO733.
US2003199436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                     ADF27868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199437-A1.
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Human secreted/transmembrane protein, PRO733.
US2003211091-A1.
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                              ADF33765 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003194780-A1.
                                                                                                                                            Human PRO polypeptide #205.
192003199029-A1.
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US2003199063-A1.
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23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; SC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                            16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
RESULT 234
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RESULT 232
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Best Local Similarity
RESULT 231
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ID AD
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PN US
PD 23
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Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207360-AI.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE91747 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003199058-AI.
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Novel human secreted and transmembrane protein PRO733
                                                                                                                                                                                                                      (GETH) GENENTECH INC.
ry Match
F. Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                      Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
RESULT 240
ID ADF26608 standard; protein; 229 AA.
DE Human secreted/transmembrane protein, PRO733.
PN US2003199674-A1.
                                                                                                                                                     ADF34397 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003194410-A1.
                                                                                                                                                                                                                                                                                                 ADF46634 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195344-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 229 AA
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US2003207376-A1.
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                                                US20031550.
23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%;
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RESULT 248
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ry Match 99.5%;
t Local Similarity 99.6%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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Human PRO polypeptide #205.
US2003207352-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                 Best Local Similarity RESULT 241
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Best Local Similarity
RESULT 249
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ADF98088 standard;
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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.
(ery Match 99.5%; Score 1171; DB 8; Length 229;
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Novel human secreted and transmembrane protein PRO733
US2003207424-A1.
                                                                                                                                                                                     Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
ADF98659 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003208055-A1.
                                                                                                             ADG03490 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207351-A1.
                                                                                                                                                                                                                             ADF99211 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003207353-A1.
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US2003207359-A1.
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Human PRO polypeptide #205.
US2003207425-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG15586 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003219885-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF96984 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                          US2003220
06-NOV-2003.
(GETH ) GENENTECH INC.
11-1-1-1 SP.5%; SC
                                                                                                                                               USACVC-2003.
06-NOV-2003.
(GETH ) GENENTECH INC.
99.54;
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US2003207357-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                    06-NOV-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 252
                                                                                 Best Local Similarity RESULT 250
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Best Local Similarity
RESULT 256
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Best Local Similarity
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Best Local Similarity
RESULT 257
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RESULT 254
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                                                                      Query Match
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Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG24943 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207427-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG07240 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207350-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AUSOVI/92 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207356-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG62055 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                   Novel human secreted and transmembrane protein PRO733 US2003207389-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG60951 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207390-A1.
                                                                      Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                             Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

99.5%; Score 1171; DB 8;

ery Match

99.5%; Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                              ADG06169 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207374-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG04042 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                    protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG55287 standard; protein; 229 AA
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06-NOV-2003,
(GETH ) GENENTECH INC.
99.5%;
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99.6%;
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(GETH) GENENTECH INC.
Query Match: 99.5%;
Best Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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1 (GETH) GENENTECH INC.
10-CHY Match 99.5%;
Best Local Similarity 99.6%;
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RESULT 266
Human PRO polypeptide #205.
US2003207371-A1.
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US2003207423-A1.
                                    06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                         Query Match
Best Local Similarity
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RESULT 260
                                                                                                              RESULT 259
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Length 229;

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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO733. US2003207419-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG52455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207414-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG54183 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207416-A1.
                                         Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003215905-Al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG51868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003215908-A1.
                                                                                      ADG50620 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003207803-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                  Human PRO polypeptide #205.
US2003207805-Al.
                                                                                                                                                                                                                                                                                                                                         ADH30666 standard; protein; 229 AA
Human PRO polypeptide #205.
US2003077723-A1.
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20-NOV-2003.
(GETH ) GENENTECH INC.
Werch 99.5%; SC
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06-NOV-2003.
(GETH ) GENENTECH INC.
%atch ....ity 99.5%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
Match Tritty 99.6%;
            PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 284
                                                                                                                                                                            Query Match
Best Local Similarity
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Best Local Similarity
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          06-NOV-2003
(GETH ) GENI
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PA (GETH) GENENTECH INC.

Query Match 99.5%; Score 1171; DB 8; Length 229;

Best Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                          99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                             Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207415-A1.
                                                                                                                                                                                                                                                                                                                                                                               ADG56943 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG58599 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207368-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG70965 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207420-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207363-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207421-A1.
                                                                                                                                                                                                                                            ADG57495 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003207362-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207365-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                  Human PRO polypeptide #205.
US2003207358-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                         06-NOV-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 276
ID ADG71517 st
DE Novel
-2003.
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-21 ) GENENTECH 1.
-2T MATCh
Best Local Similarity PRESULT 268
ID ADGR2256 stand**
DE Human PRO TO PN US2003.*
PD 06-TO PN PS2003.*
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                              Local Similarity
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Best Local Similarity
RESULT 273
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                                                                                                                                                                                               Query Match
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Length 229;

Best Loca RESULT 270

Query

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99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                            Novel human secreted and transmembrane protein PRO733 US2003207366-A1.
                                                                                                                                                                                                                                                                                                       ADH12657 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207378-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG59775 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207369-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel human secreted and transmembrane protein PRO733 US2003207429-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207367-A1.
                                                                                                                                                                                                                                                         99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 8;
                                                                                                                             99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG48748 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003216560-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted/transmembrane protein, PRO733
US2004005312-A1.
                                                                                                                                                                           ADG56391 standard; protein; 229 AA.
                                       ADG81152 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194793-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #205.
USZO0302231-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC.
99.5%; Sr
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                           (GETH ) GENENTECH INC
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Best Local Similarity
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Best Local Similarity
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Query Match
Best Local Similarity
RESULT 286
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RESULT 294
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ADH25669 standard; protein; 229 AA.
Human neurotrimin homologue related protein sequence SEQ ID NO:447.
EP1386931-A1.
                                                                                                                                   Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2004009548-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADII5413 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207382-A1.
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Novel human secreted and transmembrane protein PRO733
US2004009547-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207383-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO733.
US2003207349-Al.
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(GETH ) GENENTECH INC.

Sty Match 99.5%; Score 1171; DB 8;

Train eimilarity 99.6%; Pred. No. 4.1e-114;
   99.6%; Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                      ADG62644 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004006219-A1.
08-JAN-2004.
(BCTH ) GENENTECH INC.
99.5%; Score 1171; DB
st Local Similarity 99.6%; Pred. No. 4.1e-1
                     LT 295
ADG59188 standard; protein; 229 AA.
Man secreted/transmembrane protein, PRO733
US2004005657-A1.
                                                                                                                                                                                                                                                                                                                                           ADI81199 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207361-A1.
UG-NOV-2003.
                                                                        US200405.
08-JAN-2004.
(GETH ) GENENTECH INC.
Watch 99.5%; S'
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06-NOV-2003.
(GETH ) GENENTECH INC.
99.5%; S
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Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 295
                                                                                                                                                 Best Local Similarity RESULT 296
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RESULT 298
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Best Local Similarity
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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                              99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
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Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                      Length 229;
nuveled standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2004039164 A1.
26-FEB-2004.
(GETH) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                       99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                      Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL07280 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004063921-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM17446 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004048332-A1.
                                                                                                                                                       ADJ77516 standard; protein; 229 AA. Human PRO polypeptide #205. US2004038336-A1. C5-FEB-2004 (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                ADJ65638 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004038335-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM27774 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004048333-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM42498 standard; protein; 229 AA. Human PRO polypeptide #205.
US2004058424-A1.
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Human PRO polypeptide #205.
US2003077659-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                     Best Local Similarity
RESULT 305
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Best Local Similarity
RESULT 307
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RESULT 312
RESULT 304
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(HINZ/) HINZMANN B.
(HINZ/) HINZMANN B.
(HINZ/) HEIDEN CASTANOS-VELEZ E.
(CAST/) HEIDEN CASTANOS-VELEZ E.
Query Match
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
                                                                                   Length 229;
                                                                                                                                                                                                               Length 229;
                                                                                                                                                                                                                                                                                                                                            Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                          Novel bronchial cancer-associated human protein SeqID899.
DE10316701-A1.
ID AD196394 standard; protein; 229 AA.

DE Novel human secreted and transmembrane protein PR0733.

PN US2003207354-A1.

PD 06-NOV-2003.

PA (GETH ) GENENTECH INC.

QUESTY MATCH

PASEL Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 314
                                                                                                                              ADS32346 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2004203125-A1.
                                                                                                                                                                                                                                                                                                       28-OCT-2004.
(GETH ) GENENTECH INC.
Query Match
(CATH ) Generation 99.5%; Score 1171; DB 8;
Query Match
                                                                                                                                                                                                               Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                            ADT03330 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004214269-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADU50402 standard; protein; 229 AA PR0733, SEQ ID 447. US2004223964-A1.
                                                                                                                                                                                                                                                                                                                                                                                            ADU06673 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 229 AA
                                                                                                                                                                  US20044...
14-0CT-2004.
(GETH ) GENENTECH INC.
99.5%;
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Local Similarity 99.6%;
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US2005014226-A1.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GERRITSEN M E.
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GURNEY A L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WILLIAMS P M.
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STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BOTSTEIN D.
DESNOYERS L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LUAVIN I J
                                                                                                                                                                                                                        Best Local Similarity
RESULT 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KUO S S.
NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EATON D L.
FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAN J.
PAONI N F.
ROY M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADW49681 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAO W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TUMAS
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ASHK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TUMA/)
(WILL/)
(WOOD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DESN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FERR/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FONG/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GODD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GURN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PAON/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KUOS/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAPI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SHEL/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GODO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PANJ
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FILVAROFF E.

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AD MED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEB14127 standard; protein; 229 AA.
Cancer cell diagnosis method-related human protein - SEQ ID 410.
US2005153396-A1.
                                                                                                                                                                                           OT-APR-2005.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 9; Length 229;
ery Match 99.6%; Pred. No. 4.1e-114;
 Length 229;
                                                                                                                Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99.5%; Score 1171; DB 9; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                           MOZUCE 2005.
(GETH ) GENENTECH INC.
(GETY MAtch 99.5%; Score 1171; DB 9;
lery Match 99.6%; Pred. No. 4.1e-114;
 99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                   ADZ52342 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2005084935-A1.
                                                                                                                                                    ADZ03381 standard; protein; 229 AA.
Human secreted/transmembrane PRO733 protein.
US2005074837-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEA99614 standard; protein; 229 AA.
Human PRO733 protein SEQ ID NO:2.
US2005118635-A1.
                                           protein; 229 AA
                                      ADVI4737 standard; protein; 22
PRO polypeptide SEQ ID NO 543.
WO2005016962-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUN-2005.

(GODD/) GODDARD A.

(GORN/) GODWSKI P J.

(SMIT/) SMITH V.

(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                          ASHKENAZI A J.
BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PAN J.
PAONI N F.
ROY M A.
SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WILL/) TUMAS D.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                    BOTSTEIN D.
DESNOYERS L.
EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GODOWSKI P J.
GRIMALDI J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BAKE/) BAKER K P.
(BERE/) BERESINI M.
(DEFO/) DEFORGE L.
(DESN/) DESNOYERS L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KUO S S.
NAPIER M A.
                                                                                                                                                                                                                          Query Match
Best Local Similarity
Query Match
Best Local Similarity
RESULT 319
                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                              FERRARA N.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 322
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Best Local Similarity
RESULT 323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GURNEY A L.
HILLAN K J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                         GAO W.
                                                                                                                                                                                                                                                                                                                                       BOTS/)
                                                                                                                                                                                                                                                                                                                                                                                                FERR/
                                                                                                                                         RESULT 320
                                                                                                                                                                                                                                                     RESULT 321
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Human cancer-related PRO protein amino acid sequence - SEQ ID 410.
US2006040351-A1.
                                                                                                                                                                                                      Best Local Similarity 99.6%; Pred. No. 4.1e-114; ESSUL 229; RESULT 324

ID AED48170 standard; protein; 229 AA.

DE Human PRO733 amino acid sequence.

PN US2005227342-A1.

PD 13-0CT-2005.

PA (GETH ) GENENTECH INC.

QUETY MATCh

QUETY MATCh

RESULT 329;

RESULT 329

DE Human PRO amino acid sequence, seq id 410.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 99.5%; Score 1171; DB 10; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-4PR-2006.
(GETH ) GENENTECH INC.
(ery Match 99.5%; Score 1171; DB 10; Length 229;
ery Match 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 1171; DB 10; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99.5%; Score 1171; DB 9; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                         Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEH18401 standard; protein; 229 AA.
Human PRO protein amino acid sequence - SEQ ID 447.
US2006078964-A1.
                                                                                                                                                                                           99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 99.6%; Fred. No. 4
RESULT 326
ID AEGS8272 standard; protein; 229 AA.
DE Human PRO733 polypeptide SEQ ID No: 410.
PD 06-APR-2006.
PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEI43917 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEI17335 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99.5%;
                              (GDDO/) GODOWSKI P J.
(GURN/) GURNEY A L.
(SHEK) SHERWOOD S.
(SMIT/) SMITH V.
(STEW/) STEWART T A.
(TUNA/) TUNAS D.
(WATA/) WATANABE C K.
(WADD/) WOOD W I.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                       03-NOV-2005.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAO W.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SMITH V.
STEWART T A.
TUMAS D.
WATANABE C K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GODDARD A. GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BAKER K P.
BERESINI M.
DEFORGE L.
DESNOYERS L.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GURNEY A L. SHERWOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 329
                  GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZHANG Z.
                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BERE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DESN/)
(FILV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GAOW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SMIT/)
(STEW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ZHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DEFO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GODD/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WATA/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GURN)
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AEI17411 standard; protein; 229 AA.
Human cancer treatment biomarker amino acid sequence - SEQ ID 298.
US2006121511-A1.
Human cancer treatment biomarker amino acid sequence - SEQ ID 222.
                                                                                                                  Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 229;
                                                                                                                                                                                                                                                                                                                 Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99.0%; Score 1165; DB 7; Length 229; 99.1%; Pred. No. 1.8e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             wuery Match 70.9%; Score 835; DB 7; Length 193;
Best Local Similarity 93.1%; Pred. No. 6.4e-79;
RESULT 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUN-2003.
(CURA-) CURAGEN CORP.
70.3%; Score 828; DB 7; Length 162;
                                                                                                                                                                                                                                                                                                                                                                  AEI24000 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733, SEQ ID NO:410.
EP1672070-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY35962 standard; protein; 229 AA.
Extended human secreted protein sequence, SEQ ID NO. 211
WO9931236-A2.
                                                                                                                                                                                                                                                                                                                 99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                  99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99.1%; Score 1166; DB 2; 99.1%; Pred. No. 1.4e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99.1%; Score 1166; DB 8; 99.1%; Pred. No. 1.4e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADR03639 standard; protein; 456 AA.
Human T1/ST2 receptor binding protein.
US2004146501-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP19270 standard; protein; 229 AA. Human secreted polypeptide #121. US2004110339-A1. 10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 334
ID ADE95516 standard; protein; 229 AA.
DE Human NOVX4a protein.
PN W02003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE95518 standard; protein; 162 AA Human NOVX4b protein. WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG14988 standard; protein; 193 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOU3.

2003.

2003.

A.A.) CURAGEN CORF

LY Match

Best Local Similarity 99.

RESULT 335

ID ADR03639 standar

DE Human T1/STT

PD 29-7041.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human SECP-21 protein.
WO2003087300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-OCT-2003.
(INCY-) INCYTE CORP.
                                  08-JUN-2006.
(LEEH/) LEE H.
(SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                                      PA (LEEH/) LEE H.
PA (SHAW/) SHAW P.
PA (CLAR/) CLARK E A.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match
Best Local Similarity
RESULT 333
                                                                                                                        nest Local Similarity
                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEST ) GENSET SA.
                      US2006121511-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUN-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                    Query
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AULZ2684 standard; protein; 141 AA.
Human disease detection and treatment (MDDT) protein - SEQ ID 133.
WO2003062379-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO23233 standard; protein; 169 AA.
Human breast tumour associated protein 47-like polypeptide NOV1.
US2003027158-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 804; DB 7; Length 169;
Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 804; DB 8; Length 169;
Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                      69.8%; Score 821; DB 8; Length 162; 98.1%; Pred. No. 1.5e-77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            68.3%; Score 804; DB 4; Length 169; 97.5%; Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 57.1%; Score 672; DB 7; Length 141; Best Local Similarity 99.3%; Pred. No. 5.4e-62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 169;
                                                                                                                                           69.8%; Score 821; DB 2; Length 162; 98.1%; Pred. No. 1.5e-77;
                                        AAY35900 standard; protein; 162 AA.
Extended human secreted protein sequence, SEQ ID NO. 149.
W09931236-A2.
24-JUN-1999.
(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 804; DB 8;
Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         68.3%; Score 804; DB 6; 97.5%; Pred. No. 9.6e-76;
Best Local Similarity 98.8%; Pred. No. 2.7e-78; RESULT 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 804; DB 8;
Pred. No. 9.6e-76;
                                                                                                                                                                               RESULT 339
ID ADP19208 standard, protein; 162 AA.
DE Human secreted polypeptide #59.
PN US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                           AAB61129 standard; protein; 169 AA.
Human NOV1 protein.
WQ200075321-A2.
14-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADMS6365 standard; protein; 169 AA.
Human cell adhesion molecule NOV1.
US2003082554-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF66729 standard; protein; 169 AA.
Novel human protein NOVI.
US2003199103-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AD119766 standard; protein; 169 AA. Human NOV1 protein. US2004002134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADO60239 standard; protein; 169 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 97.5%;
RESULT 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              68.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68.3%;
97.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JUL-2003.
(INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 68.3%;
Local Similarity 97.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUL-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-OCT-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN-2004.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human NOV1 protein.
US2003134430-A1.
                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 346
ID ADL22684 standard;
                                                                                                                                                                                                                                                             (GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Length 227;

Length 133;

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RESULT 356

ID AEB81004 standard, protein; 227 AA.

DE Human putative TI/STZ receptor binding protein precursor SEQ ID NO:3.

PD 19205287606-A1.

PD 29-DEC-2005.

PA (HUMA) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEMY ) GENETICS INST INC.
52.3%; Score 615; DB 3; Length 133;
it Local Similarity 96.8%; Pred. No. 4.9e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43.0%; Score 506; DB 2; Length 120; 100.0%; Pred. No. 1.2e-44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26.9%; Score 317; DB 2; Length 61; 98.4%; Pred. No. 3.2e-25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY35901 standard; protein; 120 AA.
Extended human secreted protein sequence, SEQ ID NO. 150.
WO9911236-A2.
                                                                                                                                                                                                                AAWS4441 standard; protein; 133 AA.
Mouse novel secreted protein isolated from clone AI6_li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABB64192 standard; protein; 235 AA.
Drosophila melanogaster polypeptide SEQ ID NO 19368
WO200171042-A2.
                                                                                                                                                   52.3%; Score 615.5; DB 10; 55.0%; Pred. No. 9.3e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 416.5; DB 4;
Pred. No. 7.9e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43.0%; Score 506; DB 8; I 100.0%; Pred. No. 1.2e-44;
                                                                                                                                                                                                                                                                                                                          Score 615; DB 2;
Pred. No. 4.9e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA.) HUMAN GENOME SCI INC.

17 Match

12 Local Similarity 54.5%; Pred. No. 1.6e-55;
      Best Local Similarity 56.7%; Pred. No. 8.1e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           39.8%; Score 469; DB 5; 91.1%; Pred. No. 7.8e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP41689 standard; protein; 240 AA.
Human ovarian antigen HE2DQ62, SEQ ID NO:2821.
WO200200677-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY12211 standard; protein; 61 AA.
Human 5' EST secreted protein SEQ ID NO: 524.
WO9906554-A2.
                                                                                                                                                                                                                                                                                                                                                                                 AAB10248 standard; protein; 133 AA.
Human adult blood protein fragment AI6_1i.
WO200037630-A1.
29-UJN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP19209 standard; protein; 120 AA. Human secreted polypeptide #60. US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK35641 standard; protein; 107 AA.
Novel human polypeptide SegID7723.
WO200216439-A2.
                                                                                                        29-DEC-2005.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                    WOJEL-1998.
(GEMY ) GENETICS INST INC.
(EFY MATCh 52.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              35.4%;
                                                                                                                                                                     Best Local Similarity RESULT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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Best Local Similarity
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(GEST ) GENSET SA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEST ) GENSET.
                                                                                                                                                                                                                                                         WO9814470-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JUN-1999
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                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEE81034 standard; protein; 227 AA.
Human putative T1/ST2 receptor binding protein precursor SEQ ID NO:3.
US2005287606-A1.
                 AAU78099 standard; protein; 227 AA.
Human T1/5T2 receptor binding protein (ILIRLILG) variant sequence.
WO200216646-A1.
28-FBB-2002.
                                                                                                                                                                                                                                                                                                                                                           AAU78098 standard; protein; 227 AA.
Human TI/572 receptor binding protein (ILIRLILG) protein sequence.
WO20021646-A1.
28-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 10; Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENA-) GENAISSANCE PHARM INC.
ry Match 53.2%; Score 626.5; DB 5; Length 227;
t Local Similarity 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7; Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    53.2%; Score 626.5; DB 7; Length 227; 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                  Length 227;
                                                                                                                             Length 227;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 224;
                                                                                                                                                                                                                                                                                                DB 2;
                                                                                               (GEGA-) GENAISSANCE PHARM INC.
ry Match
L Local Similarity 56.3%; Pred. No. 9.4e-58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Occord
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
617 Match
626.5; DB 8;
617 Match
63.2%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Errynology. (REAJUL-2004. (REAJUL-2004. (REAJUL-2004. ES ASSOC BIOTECHNOLOGY. Ery Match 52.9%; Score 623; DB 8; ery Match 52.9%; Pred. No. 6.4e-57;
                                                                                                                                                                                                                                                                                              53.2%; Score 626.5; DB 2 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        53.2%; Score 626.5; DB 7 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB87501 standard; protein; 227 AA.
Human Tl receptor (TIR) ligand polypeptide.
US6586210-Bl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) 19-NOV-1996.
\ (IMMV) IMMUNEX CORP.
$2.3%; Score 616;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW09640 standard; protein; 224 AA.
Murine cytokine, ST2 ligand polypeptide
US5576191-A.
                                                                                                                                                                                    AAW00639 standard; protein; 227 AA.
Human cytokine, ST2 ligand polypeptide.
US5576191-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADN35149 standard; protein; 227 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADQ6619 standard; protein; 125 AA.
Novel human protein sequence #1592.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADI34090 standard; protein; 227 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Tl receptor ligand.
US2003100048-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human TlR ligand protein.
US2003198618-A1.
                                                                                                                                                                                                                                                     19-NOV-1996.
(IMMV ) IMMUNEX CORP.
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                                                                                                                                        Best Local Similarity
RESULT 348
                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NIJJ/) NI J.
(GENT/) GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUL-2003
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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RESULT 347
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Length 240;

DB 5;

Length 107;

Length

Length 120;

RESULT 374

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                             Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                        22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                            Length 217;
                                                                                                                                                                                             Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABJ37053 standard; protein; 217 AA.
Human breast cancer / ovarian cancer related protein #29.
WO2003000012-A2.
                                                                                                    Novel human secreted and transmembrane protein PRO1078 US2003032127-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABU89888 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003036147-A1.
                                                                                                                                                                                                                                                                                                                                   Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003027278-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted polypeptide PRO1078, SEQ ID NO:124. US:2003036159-Al.
                                                                                                                                                                                             Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027264 A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                        . Human secreted/transmembrane protein (PRO) #62.
US2003032112-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein (PRO) #62.
US2003040070-A1.
27-FEB-2003.
ABUS8461 standard; protein; 217 AA. Human PRO polypeptide #62. US2003027272-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU82767 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032113-A1.
                                                                                                                                                                                                                                            protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU96190 standard; protein; 217 AA
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                                                                                                                                                                                             22.6%;
31.6%;
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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                                                                            Query Match
Best Local Similarity
RESULT 375
                                                                                                                                                                                                        Best Local Similarity RESULT 376
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                                                                                                                                                                                                                                              ABU84324 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
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DB 6; Length 217;

23

us-10-692-730-2.spdi.rag

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU85024 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR98803 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR92226 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036160-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR78288 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054474-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR99413 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040063-A1.
                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                             Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                      ABU86254 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036146-A1.
                                                                                                                                                                                                            ABU67467 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036162-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO16326 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003027267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO18867 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044925-A1.
                                                                                                                                                                                                                                                                                                                                                      protein; 217 AA
                                                                                                      US200302-
20-FEB-2003.
(GETH ) GENENTECH INC.
- Watch 22.6%; Sr
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20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
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20-FEB-2003.
(GETH ) GENENTECH INC.
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31.6%;
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31.6%;
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Human PRO protein #62.
US2003036137-Al.
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Best Local Similarity
                 Query Match
Best Local Similarity
RESULT 395
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     20-FEB-2003.
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                                                     Length 217;
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Pred. No. 4.1e-19;
Novel human secreted and transmembrane protein PRO1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABU98799 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003013153-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU98014 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003107544-A1.
23-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU91720 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003027277-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR94666 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044926-A1.
                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                            Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                    Human secreted/transmembrane protein (PRO) #62.
US2003036149-A1.
20-FEB-2003.
                                                                                                                                                                                                                                         Human secreted/transmembrane protein (PRO) #62.
US2003044923-A1.
06-MAR-2003.
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US2003040062-A1.
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Pred. No. 4.1e-19;
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US2003036141-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU85639 standard, protein, 217 AA.
Human PRO polypeptide #62.
US2003036140-A1.
                                                                                                        protein; 217 AA
                                                                                                                                                                                                                              protein; 217 AA
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31.6%;
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 391
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Best Local Similarity
                                                                Best Local Similarity RESULT 385
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Best Local Similarity
RESULT 394
                                                                                                      ABU92621 standard;
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Best Local Similarity
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Best Local Similarity
RESULT 393
                US2003036144-A1.
                                  20-FEB-2003.
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RESULT 390

A B B B B B

SKED

RESULT 392

SEE

DB 6; Length 217;

Length 217;

9

Length 217;

DB 6;

RESULT 388

RESULT

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Length 217;

DB 6;

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RESULT 422
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                                       22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                         Length 217;
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                                                                 Novel human secreted and transmembrane protein PRO1078.
US2003022101-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO06210 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003022294-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Movel human secreted and transmembrane protein PRO1078 US200305118-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU8B714 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003036133-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR59246 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027275-A1.
                                                                                                                                                                         DB 6;
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Human secreted/transmembrane protein (PRO) #62.
US2003036123-A1.
20-FBB-2003.
                                                                                                                                                                         22.6%; Score 266.5; DB 6
31.6%; Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003027324-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                     ABO02140 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040054-A1.
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                                                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #62
US2003036124-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABU83409 standard; protein; 217 AA.
                                                                                                                                                                                                                               ABO11495 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                             22.6%;
31.6%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
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US2003032114-A1.
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR67113 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027266-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                       ABR66808 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003035148-A1.
20-PEB-2003.
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU65630 standard; protein; 217 AA.
Human secreted/transmembrane protein, SEQ ID 124
US200303036156-A1.
20-FEB-2003.
                                                                                                                                                                            Human secreted/transmembrane protein (PRO) #62. US2002040060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO03665 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036128-A1.
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Human secreted/transmembrane protein (PRO) #62.
US203044916-A1.
06-MAR-2003.
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Human secreted/transmembrane protein (PRO) #62.
US2003054483-A1.
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Human secreted/transmembrane protein, PRO1078.
US2003022298-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AB007478 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032117-A1.
13-FBB-2003.
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Human PRO polypeptide #62.
US2003332102-A1.
13-FEB-2003.
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(GETH ) GENENTECH INC.
"ortch 22.6%; Sr
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31.6%;
Query Match
Best Local Similarity
RESULT 415
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Length 217;
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Pred. No. 4.1e-19;
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           ABU95270 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003036117-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                DB 6;
                                                                                                                                                                                                                                                                                                                                         ABR70024 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003032138-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036132-A1.
20-FEB-2003.
                                                                          22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032137-Al.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR67832 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027269-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003027268-A1.
                                                                                                                                                                                                                                                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                    ABU71173 standard; protein; 217 AA. Human PRO1078 protein. US200305143-A1. 20-FEB-2003.
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                                                                                                                                                                                                                                  protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AB001498 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003008353-A1.
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31.6%;
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PA (GETH ) GENENTECH INC.
QUery Match 22.6%;
Best Local Similarity 31.6%;
RESULT 429
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31.6%;
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PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;

RESULT 431
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31.6%;
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Human PRO polypeptide #62.
US2003032130-Al.
13-FEB-2003.
                           Lery Match
Best Local Similarity
RESULT 426
ID ABU71173 stand
PN US200
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Best Local Similarity
RESULT 430
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Best Local Similarity
RESULT 428
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Best Local Similarity
RESULT 433
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Best Local Similarity
RESULT 434
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Best Local Similarity
RESULT 432
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RESULT 425
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DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABU94960 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
1S-2003032123-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU90508 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003032108-A1.
13-FEB-2003.
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Novel human secreted and transmembrane protein PRO1078
US2003032119-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003027263-A1.
ABR68442 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027274-A1.
                                                                                                                                                Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003032135-A1.
                                                                                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                           ABU89024 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003022297-A1.
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Human secreted/transmembrane protein (PRO) #62
US2003032111-A1.
13-FEB-2003.
                                                                                                                               ABR71854 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                protein; 217 AA.
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                                                                          22.6%;
31.6%;
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Local Similarity 31.6%;
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Best Local Similarity 31.6%;
RESULT 437
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US2003022295-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                        Query Match
Best Local Similarity
RESULT 436
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Best Local Similarity
RESULT 438
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Best Local Similarity
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6; Length 217;

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Best Local Similarity
RESULT 456
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30-JAN-2003.
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                                                                                                                                                                                                                                                        Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                               Length 217;
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Novel human secreted and transmembrane protein PRO1078.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                      ABU85944 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003022300-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU82231 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003036136-A1.
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Novel human secreted and transmembrane protein PRO1078
                                                Score 266.5; DB 6;
Pred. No. 4.1e-19;
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9
Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003027271-A1. 06-FEB-2003.
                                                                                                                                                                                  Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                     Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                               Human secreted/transmembrane protein; 217 AA. US2003036125-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU83714 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032109-A1.
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27-PPB.
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US2003027280-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABU87242 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003036138-A1.
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Best Local Similarity
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RESULT 455
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                                                            Best Local Similarity RESULT 446
                                                                                                                                                                   Local Similarity
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13-FEB-2003.
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                                                                                                                                                                                  RESULT 447
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                              22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Novel human secreted and transmembrane protein PRO1078.
US2003032131-A1.
                                                                                                                                                                                   ABU93980 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003036155-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                      ABU99813 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      O6-FEB-2003.
(GETH ) GENENTECH INC.
22ry Match 22.6%; Score 266.5; DB 6;
                                                 ABRS9792 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032120-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR66503 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027281-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR90921 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US2003040058-A1.
27-FRB-2003.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
22.6%; Score 266.5; DB 6, 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003032129-A1.
13-PEB-2003.
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US2003017540-Al.
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US2003032106-A1.
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US2003032103-A1.
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RESULT 461
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Best Local Similarity
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DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human secreted and transmembrane protein PRO1078. US2003036154-Al. 20-FBB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                            Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                            ABR70329 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032139-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                          ABR65893 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036165-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6;
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                                                                                                                                                                                                                                                                                                                                                                 DB 6;
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Pred. No. 4.1e-19;
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                                                       Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                       Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein (PRO) #62
US2003036142-A1.
ABO04580 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032107-A1.
13-FEB-2003.
                                                                                                                                                                                              protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #62.
US2003032110-Al.
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US2003036145-Al.
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20-FEB-2003.
(GETH ) GENENTECH INC.
12.6%; S/
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31.6%;
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US2003022301-A1.
30-JAN-2003.
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Best Local Similarity
RESULT 469
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Best Local Similarity
RESULT 470
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RESULT 474
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Best Local Similarity
RESULT 468
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Best Local Similarity
RESULT 471
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Best Local Similarity
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                                                                  Best Local Similarity
                                                                                                                                                                                             ABU98494 standard;
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                                                      Query Match
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                                                                                 RESULT 467
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032122-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6;
                                                 DB 6;
                                                                                                                                                                                                               ABR70939 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040069-A1.
27-FEB-2003.
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Pred. No. 4.1e-19;
                                               Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
Human secreted/transmembrane protein (PRO) #62.
US2003044931-A1.
06-MAR-2003.
                                                                                    Human secreted transmembrane protein (PRO) #62. US2003036150-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU84629 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003032116-A1.
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US2003036152-Al.
20-FEB-2003.
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Human secreted/transmembrane protein (PRO) #62
US2003017543-A1.
                                                                                                                                                                                                                                                                                                                               ABU87547 standard; protein; 217 AA
Human PRO polypeptide #62.
US2003022293-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 483
ID ABU80096 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                217 AA
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31.6%;
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31.6%;
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Human PRO polypeptide #62.
US2003032128-A1.
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Human PRO polypeptide #62.
US2003017541-A1.
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US2003036139-A1.
20-FEB-2003.
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                                                                Best Local Similarity
                                                                                                                                                                                Best Local Similarity
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Best Local Similarity
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                                                 Query Match
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13-MAR-2003
(GETH ) GEN
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 22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                              22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                            ABU96789 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003032140-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO04985 standard, protein, 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003008352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR80878 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049741-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR70634 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003040076-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR73989 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036135-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054455-A1.
                                                                                                                                                                                                                                                                                                                                                                                                     22.6%; Score 266.5; DB·6; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003044922-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #62
US2003032118-Al.
                                                      ABUI0571 standard; protein; 217 AA.
Human secreted/transmembrane protein #62.
US2002127584-A1.
                                                                                                                                                                                                ABU95580 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032115-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO05600 standard; protein; 217 AA
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27-FEB-2003.
(GETH ) GENENTECH INC.
"arch '2.6%; SC
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09-0AN-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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31.6%;
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
RESULT 490

ID ABR70634 standard; p:
DE Human secreted polype
PN US2003040076-A1.
PD 27-FEB-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 491
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RESULT 492
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RESULT 495
Query Match
Best Local Similarity
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                                     RESULT 487
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PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 504
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PA (GETH ) GENENTECH INC.

Query Match 22.6*; Score 266.5; DB 6; Length 217;

Best Local Similarity 31.6*; Pred. No. 4.1e-19;

RESULT 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 217;
                                                                                                                                                                Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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               Length 217;
                                                                                                                                                                                                       ADEMOUBTY Standard; protein; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                           ABR88481 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068743-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM77302 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054479-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM07948 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068752-A1.
                                                    Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6;
Pred. No. 4.1e-19;
                 Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO28786 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068685-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO11531 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068725-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO40428 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003068682-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO35853 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068701-A1.
10-APR-2003.
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10-APR-2003.
(GETH ) GENENTECH INC.
Match 11-11-11 22 6%;
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20-MMR-2003.
(GETH ) GENENTECH INC.
"Match "lirity 31.6%;
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RESULT 498
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(GETH ) GENENTECH INC.
22.6%;
             Query Match 22.6%;
Best Local Similarity 31.6%;
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                                                                                                                            13-MAR-2003.
(GETH ) GENENTECH INC.
GENENTECH INC.
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Best Local Similarity
RESULT 500
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A D B E D
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                                       Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                              Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003104539-AL.
                                                                                                                                        Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM17225 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054459-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR95276 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040071-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                   DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADMINISTRANGARD, PROTEIN; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003040075-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR94971 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044930-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                             Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                        Human secreted/transmembrane protein (PRO) #62
US2003073180-A1.
                                                                                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein (PRO) #62.
US2003036131-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO21514 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054471-A1.
                                                                                                                                                                             ADA77876 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                      217 AA
                                                                           protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR97778 standard; protein; 217 AA
                                                  JULY 206

JULY 206

JULY 206

JULY 206

Human PRO polypeptide #62.

JULY 31.6%; PARLON 2003

JULY BENENTECH INC.

March 2003

JULY 31.6%; Py
                                                                                                                                                                                                   US20030-2003.
27-FEB-2003.
(GETH ) GENENFECH INC.
22.6%; S/
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20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; /
(GETH ) GENENTECH INC. 22.6%;
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31.6%;
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31.6%;
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                                             Best Local Similarity RESULT 506
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RESULT 515
ID ABR97778 standard: 1
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RESULT 508
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Best Local Similarity
RESULT 513
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Best Local Similarity
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RESULT 509
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Best Local Similarity
RESULT 511
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                                                                                                                                                                                                                                                                                                                                                                      ABO03055 standard;
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                                       Query Match
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Length 217;
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                                                                        Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064452-A1.
                                                                                                                         ABR87566 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068705-A1.
                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                              ABM77607 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054473-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM06118 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM03624 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068722-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM35075 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073183-A1.
                                                                                                                                                                                                                                                                                                                                                                                            ABM27837 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064440-A1.
03-APR-2003.
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Human secreted polypeptide PRO1078, SEQ ID NO:124
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Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6
Pred, No. 4.1e-19;
                                                                       Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO48094 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
                 US200300...
03-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
                                                                                                                                                                                                                                                                                            US200303...
20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
                                                                                                                                                                 PD 10-ARR-2003.
PA (GETH ) GENENTECH INC.
QUETY MATCH 22.6%;
Best Local Similarity 31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                     Query Match
Best Local Similarity
RESULT 516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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RESULT 518
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RESULT 523
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Best Local
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RESULT
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Length 217;

Best Loca RESULT 526

RESULT 525

RESULT 528

A B B B B B

BEER

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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(GETH) GENENTECH INC.

(GETH) GENENTECH INC.

22.6%; Score 266.5; DB 6;

ery Match

21.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR90616 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM76083 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104548-A1.
                22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM76387 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124 US203082717-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                   ABO41343 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US20031068695-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AB002445 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO03360 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036127-A1.
20-FBB-2003.
                                                                                                                                                                                                          ABO36158 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068703-A1.
                                                                                                                                                                                                                                                                                                                                                 AB043687 standard; protein; 217 AA
Human PRO polypeptide #62.
US2003068732-Al.
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
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                Query Match
Best Local Similarity
RESULT 534
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Best Local Similarity
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                                                                                                                        10-APR-2003
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                                  Length 217;
                                                                                                                                                                                                                                                                                                                      DB 6; Length 217;
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                                                                                                                                                                           DB 6; Length 217;
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                                22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM29057 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068721-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068707-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM09473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073175-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PRO1078, SEQ ID NO:124 022003064463-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM07033 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068699-A1.
                                                                                                                                                                                                                            ABM11608 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064447-A1.
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                                                                                                                                                                                                                                                                                                                                                                ABMO2709 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073184-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 6;
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(GETH ) GENENTECH INC.

22.6%; Score 266.5; DB 6;

ery Match 22.6%; Pred. No. 4.1e-19;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 532
                                                                                                                                                                           Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003064451-Al.
                                                                                AB024597 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003065159-A1.
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03-APR-2003.
(GETH ) GENENTECH INC.
Watch 22.6%; SC
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03-APR-2003.
(GETH ) GENENTECH INC.
"Match 17-rity 31.6%;
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31.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 531
                                                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
                                                                                                                                       03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Lery Match
Best Local Similarity
RESULT 533
ID ABM09473 stard
DE Human Ser
                  GENENTECH INC
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                                                   Best Local Similarity
 03-APR-2003
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                                    Query Match
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Length 217;

Length 217;

DB 6;

Length 217;

Length 217;

Length 217;

Length 217;

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DB 6; Length 217;
                                Length 217;
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                                22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                            22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068718-A1.
                                                                         ABR73684 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US200305468-A1.
20-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                            DB 6;
                                                                                                                                                                                                                                                                                                                                     ABR94361 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
082003044917-A1.
06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR75868 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044929-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR71244 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003059880-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR93141 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064465-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted polypeptide PRO1078, SEQ ID NO:124 US203054478-A1.
                                                                                                                                                                                                                                                                                                                                                                                                     22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                              USZUCZU.
CG-MAR-2003.
(GETH ) GENENTECH INC.
(GETY MATCh 22.6%; Score 266.5; DB 6;
(ery Match 22.6%; Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 6;
TAGRAI Similarity 31.6%; Pred. NO. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 6 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #62.
US2003054470-A1.
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Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
22.6%; Score 266.5;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                          Best Local Similarity
RESULT 545
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RESULT 550
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RESULT 551
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Best Local Similarity
RESULT 552
                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 547
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                                                 Best Local Similarity
US2003036130-A1
20-FEB-2003.
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                                Query Match
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                       Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 6; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             мьміля standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                       ABM04903 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068727-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM08863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                  10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 6;
2ry Match 22.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO36463 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068714-A1.
                                 ABO30006 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003064461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO39513 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003068776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM10388 standard; protein; 217 AA. Human secreted polypeptide PRO1078, US2003069407-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO35548 standard, protein, 217 AA.
Human PRO polypeptide #62.
US2003068758-Al.
                                                                                                                                                                                          protein; 217 AA
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10-APR-2003.
(GETH ) GENENTECH INC.
Watch 22.6%; Sr
                                                                                                                                                                                                                            US2003usc...
10-APR-2003.
(GETH ) GENENTECH INC.
22.6%;
                                                                                         PD 03-APR-2003.

PA (GETH) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
QUERY MATCh 22.6%;
Best Local Similarity 31.6%;
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Query Match
22.6%;
Best Local Similarity 31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Local Similarity 31.6%;
                                                                                                                                                                                    ABO31215 standard, protein.
Human PRO polypeptide #62.
US2003068724-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                   RESULT 553
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                        Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR86956 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM28142 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054476-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR97168 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM15268 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068692-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM10998 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049782-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                            Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 6
Pred, No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO23682 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032134-A1.
13-FEB-2003.
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Human secreted/transmembrane protein (PRO)
US2003068733-A1.
                                 ABO52059 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049768-A1.
                                                                                                                                                                                                                                                                                  ABO52364 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049771-A1.
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                                                                                         US200304...
13-MAR-2003.
(GETH ) GENENTECH INC.
400 CA. 100 CA
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31.6%;
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31.6%;
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2003.

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Lift ) GENENTECH IN STATCH

Best Local Similarity 3

RESULT 563

ID ABO52364 standa-
DE Human PRO re
PD 13-0030'
PD 13-0050'
PA
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(GETH ) GENENTECH INC.
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J32134-A1.

rEB-2003.

Lory Match
Best Local Similarity
RESULT 565
ID ABR91168 stand>
DE Human secr.
PN US2003^c
PD 20
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RESULT 569
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RESULT 564
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RESULT 570
RESULT 562
1D ABO520
DE Human
PN US2003
PD 13-MAR
PA (GETH
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ID ABM06
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13-FWAN-2005.
(GETH ) GENENTECH INC.
:ry Match
+ Tocal Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-AFK-2003.
(GETH ) GENENTECH INC.
iry Match
22.6%; Score 266.5; DB 6; Length 217;
... f.n.nl Similarity 31.6%; Pred. No. 4.1e-19;
                                                                              Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-APR-2003.
(GFTH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068709-A1.
10-APR-2003.
                                                                                                                                                                                                                                                                             Human secreted polypeptide PRO1078, SEQ ID.NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM33143 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003087374-A1.
08-MAY-2003.
                                                                                                                                    ABM04234 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM35380 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073179-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM07643 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068751-A1.
                                                                                                                                                                                                                                     Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                            Score 266.5; DB 6
Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO50229 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049777-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU99223 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US200307.
13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S'
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10-APR-2003.
(GETH ) GENENTECH INC.
*** MATCh | 132.6%;
                        /S200.5.v...
10-ARR-2003.
(GETH ) GENENTECH INC.
22.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.

ry Match 22.6%;

t Local Similarity 31.6%;
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US2003049773-A1.
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                                                                                            Best Local Similarity RESULT 572
                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
RESULT 574
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RESULT 578
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ABC41953 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049745-A1.
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13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
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31.6%;
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31.6%;
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31.6%;
                                                13-MAR-2003.

1 (GETH) GENENTECH INC.

Query Match 22 6%;

Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                       Best Local Similarity RESULT 592
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                                                                                                               RESULT 591
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                  Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
              Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                      ABM18445 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054480-A1.
                                                                                                                                                                                                                                                                                                                                        ) 20-MAR-2003.
A (GETH ) GENENTECH INC.
QUETY MATCh 22.6%; SCOTE 266.5; DB 6;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                   ABR97473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003059885-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 22.6%; Score 266.5; DB 6;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003073169-A1.
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                                                                                                                            Score 266.5; DB 6;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR80573 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049740-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM01184 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049770-A1.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                            Human secreted/transmembrane protein (PRO) #62. US2003036164-Al.
                                                                                                                                                                         ABO05905 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040074-A1.
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13-MAR-2003.
(GETH ) GENENTECH INC.
44+Ch 22.6%; Sr
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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31.6%;
              22.6%;
31.6%;
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
      Query Match
Best Local Similarity
RESULT 581
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Best Local Similarity
RESULT 583
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                                                                                                                                         Best Local Similarity
RESULT 582
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RESULT 590
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27-FEB-2003
                                                                                                                              Query Match
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RESULT 584

RESULT 585

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PN DE

DE DE

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10.4PR-2003.
(GETH ) GENENTECH INC.
(ery Match 22.6%; Score 266.5; DB 6; Length 217;
-- rocal Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1/-AFR 2005.
(GETH) GENENTECH INC.
22.6%; Score 266.5; DB 6; Length 217;
FT Local Similarity 31.6%; Pred. No. 4.1e-19;
Length 217;
                                                                                                                                                                                  Length 217;
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Pred. No. 4.1e-19;
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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 594
ID ABM32838 standard; protein; 217 AA.
DE Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                           ABMI0083 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003067478-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM02404 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059886-A1.
DB 6;
                                                                                                                                                                                      DB 6;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003087373-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM74863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096353-A1.
22-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR96253 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US2003054458-A1.
20-MAR-2003.
                                                                                                                                                                                    Score 266.5; DB 6
Pred. No. 4.1e-19;
Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                          #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                    AB038598 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068773-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA79668 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003073173-A1.
                                                             ABO42563 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003049751-A1.
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DB 6; Length 217;
                                                                                                                                                                              22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                           ABR86651 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM23872 standard; protein; 217 AA.
Human secréted polypeptide PRO1078, SEQ ID NO:124
US2003068735-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM29667 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064456-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM22262 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068753-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM22042 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068742-A1.
                                                                                                 ABR86346 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049758-A1.
                                                                                                                                                                                                                                                                                                                                                       ABM16615 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064448-A1.
                                                     22.6%; Score 266.5; DB 6 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003068693-A1.
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(GETH ) GENENTECH INC.
22.6%; Score 266.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted/transmembrane protein (PRO) US2003068756-A1.
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RESULT 606
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31.6%;
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Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Lry Match
Best Local Similarity -
RESULT 600
ID ABR86346 stand*
DE Human secr*
PN US2003*
                                                                                                                                                               (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                           Best Local Similarity RESULT 601
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                                                                                                                                                                                                                                                                                                                                        RESULT 602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 603
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Query Match
22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.

22.6%; Score 266.5; DB 6; Length 217;
t Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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                                                                                                                  Length 217;
                                                                                                                                                                                                                                                       Length 217;
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                                                                                                                                                                                                                                                                                                            ABM66396 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068737-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM75778 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104547-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM34058 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR96558 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
                                       ABM28447 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003082715-A1.
                                                                                                                                                                         ABM28752 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003082716-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM34363 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003100061-A1.
29-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                  Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein (PRO) #62. US200305454-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO22124 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054477-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO20294 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032125-A1.
13-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US20031000.
29-MAY-2003.
(GETH ) GENENTECH INC.
22.6%; S'
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10-APP-2003.
(GETH ) GENENTECH INC.
22.6%;
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OS-JUN-2003.
(GETH ) GENENTECH INC.
TW. Match '11 ATIEV 31.6%;
                                                                                                                    22.6%;
                                                                                                                                                                                                                                                       22.6%;
31.6%;
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31.6%;
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31.6%;
                                                                                                                                  Best Local Similarity RESULT 610
                                                                                                                                                                                                                                                                Best Local Similarity RESULT 611
                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match
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                                                                                                                      Ouery Match
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DB 8; Length 384;
22;
                                                                                                                                                                                                                                                          Score 82; DB 8; Length 366; Best Local Similarity 21.5%; Pred. No. 21; RESULT 1067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PARA-) PARADIGM GENETICS INC. (UTNC-) UNIV NORTH CAROLINA. 7.0%; Score 82; DB 8; Length 384; Best Local Similarity 21.5%; Pred. No. 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 7.0%; Score 82; DB 4; Length 610; Best Local Similarity 26.3%; Pred. No. 44; RESULT 1073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aucry Match 7.0%; Score 82; DB 3; Length 610; Best Local Similarity 26.3%; Pred. No. 44; RESULT 1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 4; Length 610;
                                                                                                DB 8; Length 366;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 82; DB 4; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amouass6 standard; protein; 610 AA.
Mammalian toxicological response marker protein #8.
W0200136684-A2.
Nicotiana tomentosiformis GPA1 ortholog, SEQ ID 38.
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 82; DB 3;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aury 9069 standard; protein; 610 AA. Human PRO polypeptide sequence #46. WO201068948-A2. 20-SEP-2001. (GETH ) GENENTECH INC. 7.0%; Score 82; ery Match
                                                        (PARA-) PARADIGM GENETICS INC.
(UYNC-) UNIV NORTH CAROLINA.
TY MATCH 7 0%; Score 82;
T Local Similarity 21.5%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.0%; Score 82;
21.5%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH78764 standard; protein; 384 AA.
Tobacco GPA1 ortholog, SEQ ID 44.
WO2004003146-A2.
                                                                                                                                                       ADH78760 standard; protein; 366 AA.
Tobacco GPA1 ortholog, SEQ ID 40.
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                  ADH78762 standard; protein; 384 AA.
Tobacco GPA1 ortholog, SEQ ID 42.
WO2004003146-A2.
08-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAM93654 standard; protein; 610 AA.
Human polypeptide, SEQ ID NO: 3522.
EP1130094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY66653 standard; protein; 610 AA Membrane-bound protein PRO872.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY92515 standard; protein; 610 AA
                                                                                                                                                                                                                               (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA. Query Match
                                                                                                                                                                                                                                                                                                                                                                                                               (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA.
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26.3%;
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Best Local Similarity 26.3%;
RESULT 1070
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(INCY-) INCYTE GENOMICS INC.
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(HELI-) HELIX RES INST.
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                                                                                                Ouery Match
Best Local Similarity
RESULT 1066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 1068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 1072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200020604-A2.
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                                            08-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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DB 6; Length 610;
44;
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                                                                                                                  DB 4; Length 610; 44;
                                                                                                                                                                                                                                                                                                                                        7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6; Length 610;
44;
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Pred. No. 44;
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                                                                                                                                                                                                                                                   Novel human secreted and transmembrane protein PRO872 US2003032127-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABUS9069 standard; protein; 610 AA.
Növel human secreted or transmembrane protein PRO872.
US2002132252-A1.
                                       AAB65176 standard; protein; 610 AA.
Human PRO872 (UNQ439) protein sequence SEQ ID NO:113.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR66182 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027278-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR65572 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036159-A1.
20-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 6;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU99512 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003040070-A1.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                        ABU84308 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003032112-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6;
44;
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Pred. No. 44;
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Human secreted/transmembrane protein PRO872.
US2003032023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB /
Best Local Similarity 26.3%; Pred. No. 44; RESULT 1074
                                                                                                                                                                                                                                                                                                                                                                                                                                              Luery Match 7.0%; Score 82; Best Local Similarity 26.3%; Pred. No. RESULT 1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7.0%; Score 82; 26.3%; Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 82;
Pred. No.
                                                                                                                                                                                                                                7.0%; Score 82; 26.3%; Pred. No.
                                                                                                                    Score 82;
Pred. No.
                                                                                                                                                   Human PRO polypeptide #46.
052003027272-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABUS7991 standard; protein; 610 AA.
Human PRO polypeptide #23.
US2003027163-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.0%;
                                                                                                               Best Local Similarity 26.3%;
RESULT 1075
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Best Local Similarity 26.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-SEP-2002.
(GETH ) GENENTECH INC.
                                                                   WO200073454-A1.
07-DEC-2000.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1079
                                                                                                                                                                                                                                           Best Local Similarity RESULT 1076
                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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RESULT 1080
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Best Local Similarity
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                                                                                                                                                                                                                 06-FEB-2003.
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Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 628
                                             Length 217;
                                                                                                                                                                         Length 217;
                                                                                                                                                                                                                                                                                                    Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 217;
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PD 20-MAR-2003.

PA (GETH) GENENTECH INC.

Query Match

22.6%; Score 266.5; DB 6;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;

RESULT 619
                                                                                      ABR85736 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049753-A1.
13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                      22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                    ABR99718 standard; protein; 217 AA.
Ukman secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049763-A1.
13-MAR-2003.
GETH) GENENTECH INC.
22.64; Score 266.5; DB 6; St. Local Similarity 31.64; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted polypeptide PRO1078, SEQ ID.NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003073172-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6;
                                                                                                                                                                                                                                                                                                                                                ABM00574 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
17-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                               DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO38293 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO29701 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003068700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USACOURT (GETH ) GENENTECH INC. 22.6%; Score 266.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO45593 standard, protein, 217 AA.
Human PRO polypeptide #62.
US2003073182-A1.
                                                                                                                                                                                                                                                                                                                                                                                                               22.6%;
31.6%;
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Best Local Similarity 31.6%;
RESULT 627
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(GETH ) GENENTECH INC.
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RESULT 620
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RESULT 621
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Best Local Similarity
RESULT 622
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05-JUN-2003.
(GETH ) GENENTECH INC.
lery Match 22.6%; Score 266.5; DB 6; Length 217;
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(GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Fred. No. 4.1e-19;
RESULT 636
                                                                                                                                                                                                                                        Length 217;
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Pred. No. 4.1e-19;
ABM20517 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104557-A1.
                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049744-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADMINISTRATED BY PROTEIN; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064446-A1.
                                                                                                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                              ADA81395 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003092121-A1.
                                                                                                                                                                                                                                                                                          ABO16631 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003027276-A1.
                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 631
ID ABO18257 standard; protein; 217 AA.
DE Human secreted/transmembrane protein (PRO) #62.
PN US2003044920-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO22684 standard; protein; 217 AA. Human PRO polypeptide #62. US2003027265-A1. G-FEB-2003.
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13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S'
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15-MAY-2003.
(GETH ) GENENTECH INC.
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31.6%;
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31.6%;
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(GETH) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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31.6%;
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22.6%;
31.6%;
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(GETH ) GENENTECH INC.
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            Best Local Similarity RESULT 647
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RESULT 656
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RESULT 648
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RESULT 649
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                               22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM03929 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068734-A1.
                                                                                                                                                                                           Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003064458-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003068702-A1.
                                                             Human secreted polypeptide PRO1078, SEQ ID NO:124 US:2003022121-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                        AB022481 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064460-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AB037073 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO41648 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068729-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO30311 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064464-A1.
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US2003068738-Al.
10-APR-2003.
(GETH ) GENENTECH INC.
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31.6%;
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PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 31.6%;
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31.6%;
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PA (GETH ) GENENTECH INC.

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22.6%;

Best Local Similarity 31.6%;
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PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;

RESULT 645
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Best Local Similarity 31.6%;
RESULT 643
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                 Best Local Similarity
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                                                 Best Local Similarity
US2003073171-A1.
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ID ABO284
DE Human
PN US2003
PD 03-APR
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PA (GETH ) GENENTECH INC.
QUETY MATCh
22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 644
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                ABM25092 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US20031045540-A1.
05-JUN-2003.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                            Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                         ABO47789 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049747-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein (PRO) #62.
US20039779-A1.
                                                                                                                                                                                                                    ABO47484 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049742-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO48399 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049750-A1.
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ID ABMIGS20 standard; protein; 217 AA.
DB Human secreted polypeptide PRO1078,
PN US2003040078-A1.
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US2003049766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO51754 standard; protein; 217 AA Human PRO polypeptide #62.
US2003049767-A1.
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13-MAR-2003.
(GETH ) GENENTECH INC.
Watch 22.6%; Sc
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31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Query Match
22.6%;
Best Local Similarity 31.6%;
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Local Similarity 31.6%;
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DB 6; Length 217;

DB 6; Length 217;

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                              ABM75473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                       ABM25397 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104541-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM19907 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104554-Al.
05-JUN-2003.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003032136-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR71549 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032133-A1.
                                                                                                                                                                                                                                                05-JUN-2003.
(GETH ) GENENTECH.INC.
1ry Match
1ry Match 12:6%; Score 266.5; DB 6;
1ry Match 13:6%; Pred. No. 4.1e-19;
                                                                                                                                               Score 266.5; DB 6,
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                      ABO39208 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068775-A1.
10-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA81193 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049752-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO46813 standard; protein; 217 AA.
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
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05-JUN-2003.
(GETH ) GENENTECH INC.
22.6%; S
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13-MAZ-2003.
(GETH ) GENENTECH INC.
22.6%; S
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13-MMR-2003.
(GETH ) GENENTECH INC.
22.6%;
     22.6%;
31.6%;
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31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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31.6%;
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Human PRO polypeptide #62.
US2003049765-A1.
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US2003049762-Al..
     Query Match
Best Local Similarity
RESULT 666
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Best Local Similarity
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Best Local Similarity
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RESULT 667
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                                                                          Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM16310 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064449-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM09168 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073174-A1.
                                                                                                                                                                                                                                             ABR96863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054462-A1.
                                                                                                                                                                                                                                                                                                                                                                                        ABM12218 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064445-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM24177 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064441-A1.
03-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM14658 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068696-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM04539 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM06728 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068730-A1.
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                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
ABO17952 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
                                                                                                                     ABO20904 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032132-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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03-APR-2003.
(GETH ) GENENTECH INC.
Match 17-rity 22.6%;
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03-APR-2003.
(GETH ) GENENTECH INC.
22.6%;
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31.6%;
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31.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 664
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Best Local Similarity 31.6%;
RESULT 665
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31.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 659
ID ABM12218 standard; pr
DE Human secreted polype
PD 03-ARR-2003
PA (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 663
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RESULT 660
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Best Local Similarity
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Best Local Similarity
RESULT 658
                                                                   Query Match
Best Local Similarity
RESULT 657
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Best Local Similarity
RESULT 662
                                US2003044918-A1.
                                                                                                                                                                             13-FEB-2003.
                                                    06-MAR-2003.
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RESULT 661

2822

ID DE

Query Match

DB 6; Length 217;

DB 6; Length 217;

6; Length 217;

Length 217;

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RESULT
                                                         Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                 ABR73379 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054467-A1.
Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003036129-A1.
                                                                                                                                                                                                                                        ABR84821 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040057-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR73074 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027270-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABMIB140 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054469-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR76473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044932-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                            Human secreted/transmembrane protein (PRO) #62.
US2003040053-A1.
27-FEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO25342 standard; protein; 217 AA. Human PRO polypeptide #62. US2003054463-A1.
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US2003054466-A1.
                                                                                                               protein; 217 AA
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 683
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                                                                          Best Local Similarity RESULT 676
                                                                                                                                                                                                      Local Similarity
                                                                                                               ABO06868 standard;
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Best Local 8
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                                                           Query Match
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03-APR-2003.
(GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                    Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                        Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                           Length 217;
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ABR94056 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059879-A1.
                                                                                                                                                                                                                                             Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted polypeptide PRO1078, SEQ ID NO:124, US2003068760-A1.
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM05508 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003045700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM15573 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                     ABR79963 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049738-A1.
                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO30616 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064466-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO30921 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064468-A1.
                                                                                                                                                                                                                                                                                                                                                                                           AB032910 standard, protein; 217 AA.
Human PRO polypeptide #62.
US2003064453-A1.
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03-APR-2003.
(GETH ) GENENTECH INC.
"artch 22.6%; Sc
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S'
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(GETH ) GENENTECH INC.
122.6%; S/
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03-APR-2003.
(GETH ) GENENTECH INC.
Match 'larity 31.6%;
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31.6%;
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                                                                                      Query Match
Best Local Similarity
RESULT 686
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                                                                                                                                                                                           13-MAR-2003.
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Length 217;

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                    ABM26922 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted polypeptide PRO1078, SEQ ID NO:124.
022003068763-A1.
                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                            ABR88176 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068720-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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US2003036126-A1.
                                                                                                     ADA78488 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003073181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein (PRO) #62.
US2003068689-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO49924 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO50839 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO05295 standard; protein; 217 AA
USZUCZ-
13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%;
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31.6%;
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RESULT 708
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
                                                                                                                                                         17-APR-2003.
(GETH) GENENTECH INC.
OUCY MATCH
BEST LOCAL Similarity 31.6%;
                                                                                                                                                                                                                                                                                                   10-APR-2003.

1 (GETH) GENENTECH INC.

Query Match 22 6%;

Best Local Similarity 31.6%;
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(GETH) GENENTECH INC.
Query Match 22 6%;
Best Local Similarity 31.6%;
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Local Similarity 31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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                                                               Best Local Similarity RESULT 704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO39818 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                    Query Match
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                                           10-APR-2003.
(GETH ) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 6; Length 217;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                              Length 217;
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Pred. No. 4.1e-19;
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                                                                                                                         Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 6;
Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 6;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                              DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM66701 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068688-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM19602 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104552-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6;
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                                                                                                                                                                                 10-26-2-1
10-27H ) GENENTECH INC.
(GETH ) GENENTECH INC.
(Ery Match 22.6%; Score 266.5; DB 6;
(Ery Match 21.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                             ABO42258 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049748-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
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US2003068765-Al.
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003082767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO49314 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO49619 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049775-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #62.
US2003049754-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                        ABO37988 standard; protein; 217 AA
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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13-MAR-2003.

(GETH ) GENENTECH INC.

22.6%;
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;

RESULT 699
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31.6%;
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Best Local Similarity 3.
RESULT 702
ID AB049314 standar
DE Human Secret
PN US20030.
PA
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 697
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RESULT 696
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Best Local Similarity
RESULT 701
                                US2003068698-A1.
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                                                                                    Query Match
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Length 217;

Length 217;

Length 217;

Length 217;

Length 217;

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Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                           Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
     22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                            Human secreted polypeptide PRO1078, SEQ ID NO:124, US2003073176-AL.
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(GETH) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 7;
st Local Similarity 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                            ABM05813 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADMUJO14 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
10-2003068764-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM18992 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104550-A1.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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US2003049757-A1.
                                                                                                                                                                                                                                                                                                                                  ABO34938 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068728-A1.
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Human PRO polypeptide #62.
US2003049761-A1.
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13-MAZ-2003.
(GETH ) GENENTECH INC.
22.6%;
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LT-APR-2003.
(GETH ) GENENTECH INC.
22.6%;
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RESULT 729
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(GETH ) GENENTECH INC.
22.6%;
St Local Similarity 31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                Best Local Similarity RESULT 723
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Best Local Similarity
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       Query Match
                                                                                                                                                                                                                                                                                                                      RESULT 725
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RESULT 730
                                                                                       Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                      DB 7; Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049759-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM10693 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064455-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM76692 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054465-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR89396 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073170-A1.
                                                                                                                                                                                                                                                              ABR95886 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040073-A1.
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RESULT 713
ID ABR77078 standard; protein; 217 AA.
DE Human secreted polypeptide PR01078, SEQ ID NO:124
PN US2003044927-A1.
                                                                                                                                        ABM17835 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040072-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 7;
Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 7;
ery Match 31.6%; Pred. No. 4.1e-19;
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03-8FR-2003.
(GETH ) GENENTECH INC.
(EFY Match 22.6%; Score 266.5; DB 7
                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                      ABO21819 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054475-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human secreted/transmembrane protein (PRO) #62.
US2003032124-Al.
13-FEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO24292 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO19989 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                       US200305.
20-MAR-2003.
(GETH ) GENENTECH INC.
March 22.6%; SC
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03-APR-2003.
(GETH ) GENENTECH INC.
Watch 22.6%; SC
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PA (GETH ) GENENTECH INC.
QUERY MARCH 22.6%;
Best Local Similarity 31.6%;
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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Best Local Similarity
RESULT 722
ID ABR89396 stan-
DE Human ser
PN US>-
PD
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Best Local Similarity
RESULT 719
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                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 716
                                                                                       Query Match
Best Local Similarity
RESULT 714
                                                                                                                                                                                                                                                                                                                  27-FEB-2003.
                                                                                                                                                                                               27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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N N N N

AS SE

A D D E D

A B M E D

2 2 2 C

Length 217;

Length 217;

Length 217;

Length 217;

Length 217;

PER

RESULT 732

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Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                        Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                            Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.6%; Score 266.5; DB 7; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM14353 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068686-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM09778 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073178-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM34668 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104538-A1.
05-JUN-2003.
                                                                                              ABM24482 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064444-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                    ABO29396 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US20010086897-A1.
10-APR-2003.
(GETH ) GENENTECH INC.
22.64; Score 266.5; DB 7
PRY Match
51.0631 Similarity 31.68; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #62
US2003068710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO38903 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO51144 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABO03970 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003036158-A1.
20-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                 ABO31226 standard; protein; 217 AA
                                                                                                                                                                                              22.6%;
31.6%;
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RESULT 745
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(GETH) GENENTECH INC.

Query Match

Best Local Similarity 31.6%;
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
                                        Query Match 22.6%;
Best Local Similarity 31.6%;
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Best Local Similarity 31.6%;
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31.6%;
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A (GETH) GENENTECH INC.

A (GETH) 22.6%;

Best Local Similarity 31.6%;
03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                      03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                Query Match
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                                                                              RESULT 742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 746
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                                                                                                Length 217;
                                                                                                                                                                                                                                    Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059884-A1.
              ABR89091 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036119-A1.
20-FEB-2003.
                                                                                                                                                    ABR72464 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US200336120-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                          ABR74294 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036161-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR80268 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM01489 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NQ:124
US2003059882-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR87261 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM12828 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073186-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM30582 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064443-A1.
                                                                                                                                                                                                                                  22.6%; Score 266.5; DB 7; 31.6%; Pred. No. 4.1e-19;
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                                                                                              DB 7;
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                                                                                                                                                                                                                                                                                                                                                                                             DB 7;
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PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 7,
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 7
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                              Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO18562 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044921-A1.
                                                                                                                                                                                                                                                                                                                               22.6%;
31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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RESULT 740
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Best Local Similarity
RESULT 734
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Best Local Similarity
RESULT 735
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Best Local Similarity
RESULT 733
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Best Local Similarity
RESULT 736
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Length 217;

DB 7;

Length 217;

#62.

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US2003064450-A1.
03-APR-2003.
(GETH ) GENENTECH INC.
                                                              Best Local Similarity
RESULT 761
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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                                                                                                ABO31836 standard;
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                                                     Query Match
                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                  DB 7; Length 217;
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                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABMO1794 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059883-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR90006 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073177-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR93751 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054457-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM78217 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049764-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM13133 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                      ABR77683 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040067-A1.
27-FEB-2003.
                                                                                                                                                                                                                                                     ABR70893 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054456-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064442-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                        (GETH.) GENENTECH INC.

22.6%; Score 266.5; DB 7

it Local Similarity 31.6%; Pred. No. 4.1e-19;
    31.6%; Pred. No. 4.1e-19;
                                                                                                Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein (PRO) #62.
US2003054482-Al.
                         Human PRO polypeptide #62.
US2003036151-Al.
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20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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31.6%;
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31.6%;
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RESULT 755
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RESULT 759
Best Local Similarity
RESULT 751
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RESULT 756
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RESULT 757
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Best Local Similarity
                                                                                                               Best Local Similarity
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                                                                                20-FEB-2003.
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22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                    Length 217;
                                                                                                                                                                                                                                                     Length 217;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                         Length 217;
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Pred. No. 4.1e-19;
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                                                                                                                                                                       Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                      ABM08253 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068754-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM74558 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003096351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM33753 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003096358-A1.
                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104556-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 7;
Pred. No. 4.1e-19;
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                                                                                                                                    Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                       Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein (PRO) #62. US2002068681-A1.
                                                              Human secreted/transmembrane protein (PRO) #62.
US2003068731-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABO48704 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049756-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR72769 standard; protein; 217 AA. Human secreted polypeptide PRO1078, US2003036122-A1.
                                                    protein; 217 AA
                                                                                                                                                                                                                                                                                                                                     US2003000...
10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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10-APR-2003.
(GETH ) GENENTECH INC.
** MAtch '1-rity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USZUCZ-
22-MAY-2003.
(GETH ) GENENTECH INC.
MATCh ''nrity 31.6%;
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22-MAY-2003.
(GETH ) GENENTECH INC.
22.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2005...
OS-JUN-2003.
(GETH ) GENENTECH INC.
MAtch '''rrity 31.6%;
                                                                                                                                                                                                                                                       22.6%;
31.6%;
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Length 217;

Query Match

RESULT 772

SEE

Query: Match

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Query Match

1 H Z C

Length 217;

Length 217;

DB 7;

Length 217;

Length 217;

DB 7;

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Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
                                               Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                             ADM14963 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                         ABM21432 standard; protein; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003068744-A1.
                                                                                                    ABM21737 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068741-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM33448 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096357-A1.
                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM75168 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104544-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO41018 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO36768 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)·#62.
US2003068715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AB037378 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068726-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #62.
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10-APR-2003.
(GETH ) GENENTECH INC.
7--- Match --- 12 6%;
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10-APR-2003.
(GETH ) GENENTECH INC.
Match 'Trity 31.6%;
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10-APR-2003.
(GETH ) GENENTECH INC.
-- MATCh -- 12-16%;
PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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05-JUN-2003.
(GETH ) GENENTECH INC.
match 17-rity 31.6%;
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22-MAY-2003.
(GETH ) GENENTECH INC.
Match 17-rity 31.6%;
                                                                                                                                                                                                  Query Match 22.6%;
Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
:ry Match 22.6%;
                                                                                                                                                                                                                                                                                                                    PA (CBTH.) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
RESULT 782
                                                                                                                                                             10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                           DB 7; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                              DB 7; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM76997 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054464-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM30277 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068723-A1.
                                                                                                                                                                                              ABR85126 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040065-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM17530 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044928-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM22957 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068757-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                    PA (GETH) GENENTECH INC.

Query Match 22.6%; Score 266.5; DB 7

Best Local Similarity 31.6%; Pred. No. 4.1e-19;

RESULT 771
     31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                           22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
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31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                  ABO15106 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044919-A1.
                                                                #62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted/transmembrane protein (PRO) #62. US2003040077-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO28176 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064459-A1.
                                         ABO15411 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003036121-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO17241 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US200500.
03-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S
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31.6%;
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31.6%;
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Best Local Similarity

RESULT 777

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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 775
     Best Local Similarity
                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 778
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Query Match

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Query Match

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Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 798
                                                                                                                                                                    Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 22.6%; Score 266.5; DB 7; Length 217; Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 7; Length 217;
                        Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM30887 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068771-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM32107 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068708-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM32412 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068713-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM01497 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068761-Al.
                                                                                                                                                                                                                                                                                                                                                     ABM31192 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068762-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 7;
                                                                                                                                                                                                                      ABM31802 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068680-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 7
Pred, No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein (PRO) #62.
US2003054472-A1.
                                                                             ADA82559 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049755-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             #62.
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(GETH.) GENENTECH INC.
22.6%; Score 266.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD05597 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003087376-A1.
                                                                                                                                                                    22.6%;
31.6%;
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31.6%;
                        22.6%;
31.6%;
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RESULT 795
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAR-2003.
(GETH ) GENENTECH INC.
       (GETH ) GENENTECH INC.
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                                                                                                                                                    GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
RESULT 797
                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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RESULT 790
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                                     Best Local Similarity RESULT 789
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                                                                                                                                   13-MAR-2003
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ID ADB858
DE Human
PN US2003
PD 20-MAR
PA (GETH
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Ob-NOV-zous.
(GETH ) GENENTECH INC.
iry Match
22.6%; Score 266.5; DB 7; Length 217;
iry Match
31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                           Length 217;
                                                                                                                                                                                                                                                                                                                                                                   Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 217;
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ADG02592 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
                                                                                                                                                                                                                                                                           ADF95474 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003207398-A1.
                                                                                                                                     ADG01299 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003207399-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                   ADG12289 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003207392-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL32730 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003207396-A1.
06-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM30264 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003073813-A1.
17-APR-2003.
                                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI62681 standard; protein; 217 AA.
Human apoptosis-associated protein SEQ ID 124.
WO2003058021-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO polypeptide #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE74873 standard; protein; 217 AA
                                                                                                                                                                       US200320...
06-NOV-2003.
(GETH) GENENTECH INC.
(GETH) 32.6%; Sr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (XANT-) XANTOS BIOMEDICINE AG.
                                                                                                                                                                                                                                                                                                                 (GETH ) GENENTECH INC. 22.6%;
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36-NOV-2003.
(GETH ) GENENTECH INC.
22.6%;
31.6%;
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31.6%;
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31.6%;
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Best Local Similarity 31.6%;
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                                                                                                  Best Local Similarity
RESULT 799
                                                                                                                                                                                                                                         Best Local Similarity RESULT 800
                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 806
ID ADE74261 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                              US2003207397-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-2003.
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                                                                                                                                                                                                                                                                                                                                                                   Query Match
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(GETH ) GENENTECH INC.
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                  Query Match
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RESULT 824
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                                                 Length 217;
                                                                                                                                                             Length 217;
                                                                                                                                                                                                                                                                                             Length 217;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                             Ausovassy standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
082003215912-A1.
                                                                                                          Novel human secreted and transmembrane protein PRO1078 US2003215909-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADH26054 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003068770-A1.
                                                                                                                                                                                                                                                                                                                                                        Novel human secreted and transmembrane protein PRO1078 US2003215911-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM31429 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2004048334-A1.
11-MAR-2004.
                                                 Score 266.5; DB 8;
Pred. No. 4.1e-19;
                                                                                                                                                             Score 266.5; DB 8;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                         PD 20-NOV-2003.

PA (GETH ) GENENTECH INC.

Query Match
22.6%; Score 266.5; DB 8;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8;
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(GETH ) GENENTECH INC.
(ery Match 22.6%; Score 266.5; DB B;
lery Match 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 8
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
Human secreted/transmembrane protein (PRO) #62.
US2003211574-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 266.5; DB 8
Pred. No. 4.1e-19;
                                                                                                protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                          ADG00517 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH33023 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068768-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ54762 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2004023321-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Load; protein,
Judo68768-Al.

10-APR-2003.

PA (GETH) GENENTECH INC.
QUERY MAtch
Best Local Similarity 31.6%; Pre
RESULT 814
ID AD054762 standard; protein
PRO US2004023327
PA (5-FFP)
                                                                                                                                                             22.6%;
31.6%;
                                                22.6%;
31.6%;
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RESULT 815
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 812
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
RESULT 813
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Human PRO polypeptide #62.
US2004038337-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #62.
US2003215910-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                              Query Match
Best Local Similarity
RESULT 808
                                                                                                                                                                          Best Local Similarity
RESULT 809
                                                                                            ADF96086 standard;
                             13-NOV-2003.
                                                                                                                                            20-NOV-2003.
                                                                                                                                                               Query Match
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PD 13-MAY-2004.
PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 8; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                      Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 8; Length 217; Pred. No. 4.1e-19;
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Score 266.5; DB 8; Length 217; Pred. No. 4.1e-19;
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                                                                                                               ADM36476 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2004053358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               אס אייביים און אס אייבים ארא ארן ארן ארטנים; אס ארן ארן ארטנים; אס ארטנים; אס ארטנים; אס ארטנים; אס ארטנים; אט אטרטנים; אטרטנים; אייביים; אייבים; איי
                                                                                                                                                                                                                                                                                                                                                                                                          ADM40281 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2004048335-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AED50119 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2005163766-A1.
28-UUL-2005.
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Pred. No. 4.1e-19;
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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                   Score 266.5; DB 8;
Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP24144 standard; protein; 217 AA. PRO polypeptide SEQ ID NO:1322. WO2004041170-A2.
                                                                                                                                                                            US20040550-
18-MAR-2004.
(GETH ) GENENTECH INC.
****Ch 22.6%; SC
   22.6%;
31.6%;
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31.6%;
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31.6%;
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Local Similarity 31.6%;
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Local Similarity 31.6%;
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Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAY-2004.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
823
                              Best Local Similarity RESULT 817
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(NUVE-) NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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Query Match
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                                                                          RESULT 835
                                                                                                                                                                                                                                                         Human tumor overexpressed cDNA protein product PRO1078 SEQ ID NO: 124
                                                                                                                                                                                                                                                                                                                                                                     ADC32808 standard; protein; 244 AA.
Human novel contig-encoded polypeptide sequence, SEQ ID NO:2890.
WO2003029271-A2.
                                                                                                                                                                                        Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 5; Length 215;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 7; Length 244;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEE81003 standard; protein; 217 AA.
Human T1 receptor-like ligand I protein SEQ ID NO:2.
US2005287606-A1.
                                                                                                     AEG88116 standard; protein; 217 AA.
Human PRO protein amino acid sequence - SEQ ID 124
US2006074227-A1.
                                                                                                                                                                                                                                                                                                                         DB 10;
                                                                                                                                                                                        DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.4%; Score 263.5; DB 6; 31.6%; Pred. No. 8.4e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.3%; Score 262.5; DB 2; 31.1%; Pred. No. 1.1e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 263.5; DB 2;
Pred. No. 8.4e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22.3%; Score 262.5; DB 1:31.1%; Pred. No. 1.1e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 22.0%; Score 259.5; DB 5
Best Local Similarity 30.9%; Pred. No. 2.2e-18;
                                                                                                                                                                                     22.6%; Score 266.5; DB 1 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                       22.6%; Score 266.5; DB 1 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.8e-19;
                                                                          31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABG72288 standard; protein; 217 AA.
Human p24 vesicle trafficking protein Hp24-1.
US2002132999-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW49099 standard; protein; 217 AA.
Human p24 vesicle trafficking protein Hp24-1.
WO9836068-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TIR-like ligand I protein. US2003129643-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW48334 standard; protein; 217 AA.
Human T1 receptor-like ligand I.
WO9807881-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABP65162 standard; protein; 215 AA.
Hypoxia-regulated protein #36.
WO200246465-A2.
                                                                                                                                                                                                                                        AEH17432 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2002.
(OXFO-) OXFORD BIOMEDICA UK LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-1998.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-DEC-2005. (HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22.4%;
31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (INCY-) INCYTE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (INCY-) INCYTE PHARM INC.
Best Local Similarity S
RESULT 826
ID AEG88116 standar
DE Human PRO PT
PN US200607
PD 06-
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ID ABP65162 stander

PD 13-
PA
                                                                                                                                                        06-APR-2006.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                  Best Local Similarity RESULT 827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                       (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                        US2006094864-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                        10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-AUG-1998
                                                                                                                                                                                                                                                                                          04-MAY-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PNE
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ADP04573 standard, protein, 208 AA.
Sea squirt protein with tissue specific expression in development Seq168.
JP2004057129-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB74750 standard; protein; 201 AA.

Human secreted protein sequence encoded by gene 18 SEQ ID NO:59.

WO2001775-A2.

22-FEB-2001.

(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB74759 standard; protein; 201 AA.
Human secreted protein sequence encoded by gene 18 SEQ ID NO:68.
WO200112775-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB74760 standard; protein; 201 AA.
Human secreted protein sequence encoded by gene 18 SEQ ID NO:69.
WO200112775-A2.
                                                                                                                                                                                                                                                                                                                                Score 236.5; DB 7; Length 201; Pred. No. 5.2e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 04-NOV-2004.
PA (FIVE-) FIVE PRIME THERAPEUTICS INC.
Query Match 19.8%; Score 233.5; DB 8; Length 552;
Best Local Similarity 30.8%; Pred. No. 4.6e-15;
RESULT 840
                                                                                                                                  20.1%; Score 236.5; DB 7; Length 201; 29.6%; Pred. No. 5.2e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AEBBB198 standard; protein; 208 AA.
Human protein highly expressed in cancer tissue SEQ ID NO 49.
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (FIVE-) FIVE PRIME THERAPEUTICS INC.

ry Match
19.9%; Score 234; DB 9; Length 208;
t Local Similarity 28.7%; Pred. No. 9.9e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19.8%; Score 232.5; DB 4; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19.8%; Score 232.5; DB 4 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 234; DB 5;
Pred. No. 9.9e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19.8%; Score 233; DB 8; 29.2%; Pred. No. 1.3e-15;
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(HUMA-), HUMAN GENOME 19.8%; Score 232.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2004.
(KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
ADD44746 standard; protein; 201 AA.
Rat Protein Q63524, SEQ ID NO 10175.
W02003016475-A2.
27-FEB-2003.
                                                                                                                                                                                                ADD44750 standard; protein; 201 AA.
Rat Protein Q63524, SEQ ID NO 10179.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                ABP69422 standard; protein; 208 AA.
Human polypeptide SEQ ID NO 1469.
WO200270539-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AD002684 standard; protein; 552 AA.
Novel human polypeptide segid 1151.
WO2004093804-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-FEB-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                  20.1%;
29.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19.9%;
28.7%;
                                                             A (GEHO ) GEN HOSPITAL CORP.
A (FARB ) BAYER AG.
Query Match
                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
RESULT 841
                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                    27-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-AUG-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2002
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RESULT 856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 858
                                                                                                  A P D S E D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADL/8591 standard; protein; 201 AA.
Albumin fusion protein related therapeutic protein X, SEQ ID No 2073.
US2004010134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL/8593 standard; protein; 201 AA.
Albumin fusion protein related therapeutic protein X, SEQ ID No 2075.
US2004010134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human heat mitochondrial protein as a therapeutic target SeqID717 WO2003087768-A2.
                                                                                                              Length 201;
                                                                                                                                                                                                                                                                                                                                                                    Length 201;
                                                                                                                                                                                                                                          Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 201;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 8; Length 201;
                                                                                       (HUMAN) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                 ABG65324 standard; protein; 201 AA.
Human albumin fusion protein #1999.
W0200177137-A1.
18-OCT-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
st Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19.8%; Score 232.5; DB 7; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 232.5; DB 7;
Pred. No. 1.4e-15;
                                                                                                                                                                                                                                         DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 7;
                                                                                                                                                                                                MOZOCT-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTVA-) HUMAN GENOME SCI INC.
(ery Match
(ery Match
(ery Match 19.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-SEP-2003.
(CHIL-) CHILDREN'S HOSPITAL MEDICAL CENT.
ery Match 19.8%; Score 232.5; DB 7;
29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 232.5; DB 7
Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19.8%; Score 232.5; DB 8 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                           ADD44752 standard; protein; 201 AA. Human Protein Q15363, SEQ ID NO 10181. W02003016475-A2. 27-FEB-2003 (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD44748 standard; protein; 201 AA.
Human Protein Q15363, SEQ ID NO 10177.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP65317 standard; protein; 201 AA.
Human coated vesicle membrane protein.
WO2003072827-A1.
                             ABG65327 standard; protein; 201 AA. Human albumin fusion protein #2002 WO200177137-A1.
                                                                                                                                                        ABG65326 standard; protein; 201 AA. Human albumin fusion protein #2001. WO200177137-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ68911 standard; protein; 201 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19.8%;
29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MITO-) MITOKOR.
(BUCK-) BUCK INST AGE RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ROSE/) ROSEN C A.
(HASE/) HASELTINE W A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 852
ID ADL78591 standard; pr
DE Albumin fusion protei
                                                                                                          Query Match
Best Local Similarity
RESULT 845
                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 846
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Best Local Similarity
RESULT 847
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 848
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Best Local Similarity
RESULT 849
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                                                                               18-OCT-200:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                 RESULT 844
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ABM81493 standard; protein; 201 AA.
Tumour-associated antigenic target (TAT) polypeptide PRO4793, SEQ:3848.
WO2004030615-A2.
                                                                                  Albumin fusion protein; 201 AA.
SS2004010134-A1.
SS-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-APR-2006.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match
Lery Match
(ery Match 19.8%; Score 232.5; DB 10; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 10; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-2004.
(GETH ) GENENTECH INC.
(ery Match 19.8%; Score 232.5; DB 8; Length 201;
ery Match 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10; Length 201;
                                                          19.8%; Score 232.5; DB 8; Length 201; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                 Score 232.5; DB 8; Length 201; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 8; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                    Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 215;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK62498 standard; protein; 215 AA.
Disease treating protein complex-derived protein #375.
                                                                                                                                                                                                                                                                                                                                                                                    Score 232.5; DB 8;
Pred. No. 1.4e-15;
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9
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WOLVELLY CENENTECH INC.

(GETH ) GENENTECH INC.

19.8%; Score 232.5; DB B;

19.7 Match Addition 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC.
19.8%; Score 232.5; DB 10
st Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19.8%; Score 232.5; DB 10.29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 229.5; DB 6
Pred. No. 3.1e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP54500 standard; protein; 201 AA.
Human PRO protein sequence SEQ ID NO:476.
WO2004039956-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEH08866 standard; protein; 201 AA.
Therapeutic protein HCFCU69, SEQ ID 2075.
US2006084794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEH08864 standard; protein; 201 AA.
Therapeutic protein HACAB58, SEQ ID 2073.
US2006084794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEH08867 standard; protein; 201 AA.
Therapeutic protein HCB3Z39, SEQ ID 2076.
US2006084794-A1.
                                                                                                                                                                                                                                                                          Antipsoriatic protein; 201 AA. W02004028479-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABRS2923 standard; protein; 215 AA.
Protein sequence #SEQ ID 711.
EP1258494-Al.
                                                                                                                                                                                                                                                                                                                          WO2004uzc..
08-APR-2004.
(GETH ) GENENTECH INC.
19.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19.5%;
29.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                             Query Match
Best Local Similarity 29.2%;
RESULT 854
                                                                            29.28;
                                                                                                                                                                                        (ROSE/) ROSEN C A. (HASE/) HASELTINE W A.
15-JAN-2004.
(ROSE/) ROSEN C A.
(HASE/) HASELTINE W A.
                                                                      Best Local Similarity RESULT 853
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-APR-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-APR-2006
                                                        Query Match
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Length 207;

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/ Match 16.4%; Score 193.5; DB 2; Length 163; Local Similarity 36.7%; Pred. No. 1.3e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 181.5; DB 4; Length 203;
Pred. No. 3.1e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-AUG-2005.
(RIVE-) FIVE PRIME THERAPEUTICS INC.
ery Match 14.1%; Score 165.5; DB 9; Length 158;
ery Match 29.1%; Pred. No. 1.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13.9%; Score 163.5; DB 7; Length 135; 32.8%; Pred. No. 1.4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AEB88199 standard; protein; 158 AA.
Human protein highly expressed in cancer tissue SEQ ID NO 50.
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                          Query Match 15.7%; Score 185; DB 4; Length 146; Best Local Similarity 34.6%; Pred. No. 8.4e-11; RESULT 873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE10062 standard; protein; 254 AA.
Novel protein-related contig polypeptide seguence #650
WO2003054152-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Abbe4154 standard; protein; 203 AA.
Drosophila melanogaster polypeptide SEQ ID NO 19854.
WO200111042-A2.
                                                                                                                                                                                                                                            адывчячы standard; protein; 146 AA.
Human membrane or secretory protein clone PSEC0228
EP1067182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC31097 standard; protein; 135 AA.
Human novel polypeptide sequence, SEQ ID NO:1179.
WO2003029271-A2.
                                 / Match 17.1%; Score 201; DB 10; Local Similarity 28.2%; Pred. No. 2.9e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15.7%; Score 185; DB 9; 34.6%; Pred. No. 8.4e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 166; DB 7;
Pred. No. 1.8e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB24228 standard; protein; 240 AA.
Human vesicle associated protein 7 SEQ ID NO:7.
WO200060082-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 166; DB 4;
Pred. No. 1.8e-08,
                                                                                 AAW70458 standard; protein; 163 AA.
Human T1-receptor ligand III splice variant 1.
WO9838311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                        ADY63253 standard; protein; 146 AA.
Human clone PSEC0228 protein, SEQ ID 258
EP1514933-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU31151 standard; protein; 254 AA.
Novel human secreted protein #1642.
WO200179449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                        03-SEP-1998.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.4%;
25.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.1%;
27.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14.1%;
27.1%;
                (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                                                                                                                   10-JAN-2001.
(HELL-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-2003.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUL-2003.
(HYSE-) HYSEQ INC.
22-JUN-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-MAR-2005
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEJ50287 standard; protein; 207 AA.
Saccharomyces cerevisiae stress-related protein (SRP) - SEQ ID 206.
US2006137043-A1.
                                                                                                                                                                                                                         DB 10; Length 203;
                                                                                                                                                                                                                                                                                                                                                                                    DB 8; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 183;
                                                      Score 229.5; DB 7; Length 215; Pred. No. 3.1e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; Length 188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17.2%; Score 202.5; DB 8; Length 188; 30.3%; Pred. No. 1.8e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADL31427 standard; protein; 188 AA.
Human protein encoded by a full length cDNA clone SeqID 3460
EP1396543-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 210;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 207;
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABBS9997 standard; protein; 208 AA.
Drosophila melanogaster polypeptide SEQ ID NO 6783
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABB58697 standard; protein; 210 AA.
Drosophila melanogaster polypeptide SEQ ID NO 2883
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18.3%; Score 215.5; DB 5; 26.4%; Pred. No. 7.3e-14;
                                                                                                  ALANVILZ standard; protein; 203 AA.
Protein aggregation-related protein SEQ ID NO:6.
WO2006091964-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 217; DB 4;
Pred. No. 6.1e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 216; DB 4;
Pred. No. 7.8e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17.2%; Score 202.5; DB 4 30.3%; Pred. No. 1.8e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17.1%; Score 201; DB 8; 28.2%; Pred. No. 2.9e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADT87228 standard; protein; 207 AA.
Yeast Stress-related protein from gene YOR016C.
WO2004092398-A2.
                                                                                                                                                                                                                       Score 221.5; DB Pred. No. 2e-14;
                                                                                                                                                                                                                                                                                                                                                                                    Score 218.5; DB
Pred. No. 4e-14;
                                                                                                                                                                                                                                                                         ADS10510 standard; protein; 201 AA.
Human therapeutic protein - SEQ ID 747.
WO2004080148-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP69421 standard; protein; 183 AA. Human polypeptide SEQ ID NO 1468. WO200270539-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAM93624 standard; protein; 188 AA.
Human polypeptide, SEQ ID NO: 3460.
EP1130094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-MAR-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18.4%;
28.7%;
                                                        19.5%;
                                                                                                                                                                                                                                                                                                                                                                                    18.6%;
28.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18.4%;
26.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                       18.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-SEP-2001.
(HELI-) HELIX RES INST.
                                                                                                                                                                                                (UYAL-) UNIV ALABAMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 865
ID ABB56697 standard; p.
DE Drosophila melanogas:
PN W020171042-A2.
PD 27-SEP-2001.
PA (PEKE) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 868
                                                                                                                                                                                                                                      Best Local Similarity RESULT 863
                  27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                          23-SEP-2004.
(NUVE-) NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 866
                                                                           Best Local Similarity
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Best Local Similarity
RESULT 867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PEKE ) PE CORP NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2002.
(HYSE-) HYSEQ INC.
EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-SEP-2001
                                                                                                                                                                                 31-AUG-2006
                                                          Query Match
                                                                                                                                                                                                                         Query Match
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Length 146;

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Length 240;
                                                                                                                               Length 240;
                                                                                                                                                                                                                                             Score 162.5; DB 4; Length 240; Pred. No. 4e-08;
                                                                                                                                                                                                                                                                                                                                                              Length 240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 136;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8; Length 225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 8; Length 225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 228;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drosophila melanogaster polypeptide SEQ ID NO 22431.
                                                                                          DC-APR-2001.
(ZYMO) ZYMOGENETICS INC.
ery Match 13.8%; Score 162.5; DB 4;
             13.8%; Score 162.5; DB 3; 22.2%; Pred. No. 4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13.7%; Score 161.5; DB 8; 29.2%; Pred. No. 2.2e-08;
                                                                                                                                                                                                                                                                                                                                                             Score 162.5; DB 5;
Pred. No. 4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADX66737 standard; protein; 225 AA.
Plant full length insert polypeptide segid 37580.
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plant full length insert polypeptide seqid 52028
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13.7%; Score 161.5; DB 8 25.2%; Pred. No. 4.6e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 156.5; DB 8 Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4;
                                                                                                                                                           13.0%; Score 153;
                                                      AAB62393 standard; protein; 240 AA.
Human type I membrane protein ztsll
WO200129223-A2.
                                                                                                                                                                                                                                                                                   ABP64862 standard; protein; 240 AA
Human protein SEQ ID 522.
WO200259260-A2.
                                                                                                                                                                                                                                                                                                                                                                                                      ADK66189 standard; protein; 136 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               225 AA
                                                                                                                                                                                                                                            13.8%; 22.2%;
                                                                                                                                                                                                                                                                                                                                                             13.8%; 22.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13.3%;
25.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADX89364 standard; protein;
(INCY-) INCYTE PHARM INC
                                                                                                                                                                                                                26-APR-2001.
(ZYMO ) ZYMOGENETICS INC.
                                                                            ...A2.
...O ) ZYMOGENETIC.
.ery Match
Best Local Similarity ?
RESULT 881
ID AAG81378 stand?
DE Human AFP r
PN WO201?
PD 26.
                                                                                                                                                                                           (HAYA/) HAYASHIZAKI Y.
(KAMI/) KAMIYA M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                    Human phX-6 protein.
US2004019006-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
             Query Match
Best Local Similarity
RESULT 880
                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KAMI/) KAMIYA M.
(SUZU/) SUZUKI T.
(HIRO/) HIROKAWA H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
RESULT 886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TANA/) TANAKA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NAKA/) NAKAO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LIUJ/) LIU J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LIU J.
                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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ABM80706 standard; protein; 214 AA.
Tumour-associated antigenic target (TAT) polypeptide PRO19615, SEQ:1822.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ70121 standard; protein; 214 AA.
Human heat mitochondrial protein as a therapeutic target SeqID1927.
W02003087768-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 203;
Best Local Similarity 20.4%; Pred. No. 3.7e-07;
RESULT 887

ID AEL96281 standard; protein; 228 AA.

DE Drosophila melanogaster survival essential protein, SEQ:1173.

PN US7135558-B1.
                                                                                                                                                                                                                                                                                                                                                                     Length 219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 203;
                                                                                                                                    Length 228;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 214;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 214;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis thaliana protein; 253 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 55396.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAG44244 standard; protein; 216 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 55397.
BP1031405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aux62494 standard; protein; 203 AA.
Disease treating protein complex-derived protein #373.
EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                    Score 146.5; DB 8;
Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7;
                                                                                                                                                                                       ADX78578 standard; protein; 219 AA.
Plant full length insert polypeptide segid 47944.
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Abovovi standard; protein; 227 AA.
Human protein having hydrophobic domain, HP03977.
WO200149728-A2.
                                                                                                                                      13.0%; Score 153; DB 11; 20.4%; Pred. No. 3.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11.5%; Score 135; DB 3; 23.7%; Pred. No. 3.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 140.5; DB 6
Pred. No. 6.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 140.5; DB 7
Pred. No. 6.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11.5%; Score 135; DB 3; 23.7%; Pred. No. 2.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 137; DB 7;
Pred. No. 1.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.6%; Score 137; DB 8; 24.0%; Pred. No. 1.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                            Protein sequence #SEQ ID 707.
EPI258494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WOZUCZ-203.
23-OCT-2003.
(MITO-) MITOKOR.
(BUCK-) BUCK INST AGE RES.
March 11.6%; S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11.9%; 26.3%;
                                                                                                                                                                                                                                                                                                                                                                     12.4%;
22.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11.9%;
26.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
                                                                                                  14-NOV-2006.
(APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                          LIU J.
ZHOU Y.
KOVALIC D K.
SCREEN S E.
TABASKA J E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004030615-A2.
                                                                                                                                                                                                                                                                                                                                                   (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                               (KOVA/)
                                                                                                                                                                                                                                                                                                                                   (TABA/)
                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 891
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a; Length 218;

a12 AA.

a12 AA.

cest Local Similarity 23.5%; Pred. No. 4.7e-05;

RESULT 902

ID ADY22460 standard; protein; 143 AA.

DE Plant full length insert polypeptide seqid 70°

PA (SIGNA) LIU J.

PA (ZHOU/) ZHOU Y.

PA (SCRE/) SCPP

PA (SCRE/) SCPP

PA (TAPP).

PA (TAPP).

PA (TAPP).
                                           Luciy match 11.4%; Score 134.5; DB 4; Length 227; Best Local Similarity 25.1%; Pred. No. 3.2e-05; RESULT 896
                                                                                                                                                                                               11.4%; Score 134.5; DB 4; Length 227; 25.1%; Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                                          11.4%; Score 134.5; DB 7; Length 227; 25.1%; Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 130; DB 3; Length 208;
Pred. No. 8.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG10323 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 8594.
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP69317 standard, protein; 218 AA.
Human lung specific protein sequence SEQ ID NO:74.
W02003102137-A2.
(DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
8
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Best Local Similarity 22.0%; Pred. No. 3.9e-05; RESULT 899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 133; DB 5;
Pred. No. 3.9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 131.5; DB 8
Pred. No. 3.4e-05;
                                                                                                                        Human full-length polypeptide sequence #1. WO200164834-A2.
                                                                                                                                                                                                                                                 ADD19280 standard; protein; 227 AA. Human secreted protein from gene 22. WO2003052377-A2. 26-JUN-2003. (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                           ADH48778 standard; protein; 203 AA. NOV26 protein sequence, SEQ ID 62. WO200268652-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEJ61737 standard; protein; 203 AA.
Human NOV26 protein, SEQ ID NO: 62.
EP1686175-A2.
                                                                                                           227 AA
12-JUL-2001.
(PROT-) PROTEGENE INC.
(SAGA) SAGAMI CHEM RES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11.0%;
25.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11.2%;
30.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 22.0%;
RESULT 900
                                                                                                           protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-AUG-2006.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                06-SEP-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                 Query Match
Best Local Similarity
RESULT 897
                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 903
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                                                                                                                                                             07-SEP-2001.
(HYSE-) HYSEQ INC.
                                                                                                           AAU27676 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                         RESULT 898
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AEC64808 standard; protein; 219 AA.
Deregulated, liver cancer-associated protein, SEQ ID NO:123
W02005085861-A2.
(ORID-) ORIDIS BIOMED FORSCHUNGS & ENTWICKLUNGS.
                                                                                              Length 212;
                                                                                                                                                                                                                                                                                                                                                                                                                10.6%; Score 125; DB 4; Length 237; 25.2%; Pred. No. 0.00034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.5%; Score 124; DB 7; Length 219; 24.9%; Pred. No. 0.00039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10.5%; Score 123; DB 7; Length 219; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 10.5%; Score 123; DB 7; Length 193; Local Similarity 25.5%; Pred. No. 0.00041;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 10.5%; Score 124; DB 9; Length 219; Local Similarity 24.6%; Pred. No. 0.00039;
                                                                                                                                                                                                                                                      Score 126; DB 7; Length 225; Pred. No. 0.00025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rice abiotic stress responsive polypeptide SEQ ID NO:5225. WO2003008540-A2.
                                                                                                                                               ADK62522 standard; protein; 225 AA.
Disease treating protein complex-derived protein #387.
EP1338608-A2.
                    AAG10322 standard; protein; 212 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 8593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.5%; Score 124; DB 7; 24.9%; Pred. No. 0.00039;
                                                                                              Score 130; DB 3;
Pred. No. 8.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD48311 standard; protein; 219 AA.
Rat Protein CAA06212, SEQ ID NO 14009.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Protein P49755, SEQ ID NO 14007. 77-Pro-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD48307 standard; protein; 219 AA.
Rat Protein CAA06212, SEQ ID NO 14005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Protein P49755, SEQ ID NO 1367. WO2003016475-A2.
                                                                                                                                                                                                                                                                                                           AA013679 standard; protein; 237 AA. Human polypeptide SEQ ID NO 27571. W0200164835-A2. 077-SEP-2001. (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                        10.7%;
20.3%;
                                                                                            Query Match 11.0%;
Best Local Similarity 25.2%;
RESULT 905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-PEB-2003.
(GEHO ) GEN HOSPITAL CORP.
(FARB ) BAYER AG.
                                                                                                                                                                                                              27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 906
                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003016475-A2.
27-FEB-2003.
                                                          EP1033405-A2.
06-SEP-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
RESULT 904
ID AAG103
DE Arabic
PN EP1033
PD 06-SE
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Length 205;

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Score 119.5; DB 3; Length 204; Pred. No. 0.001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9.7%; Score 114.5; DB 4; Length 100; 40.4%; Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 114.5; DB 8; Length 119; Pred. No. 0.0016;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9.6%; Score 113.5; DB 7; Length 203; 23.8%; Pred. No. 0.0044;
                                                                                                                                                                                                                 DB 3; Length 225;
                                                                                                                                                                                                                                                                                                                                Length 207;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 214;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 224;
                                                AAG16692 standard; protein; 205 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 17436.
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                   AAG16691 standard; protein; 225 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 17435.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                          AAGO7344 standard; protein; 214 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4462.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAG07343 standard; protein; 224 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4461.
06-SEP-2000.
                                                                                                          10.2%; Score 119.5; DB 3; 27.3%; Pred. No. 0.001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 119; DB 3;
Pred. No. 0.0013;
                                                                                                                                                                                                                                                                                                                                10.1%; Score 119; DB 2; 20.6%; Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 119; DB 3;
Pred. No. 0.0013;
                                                                                                                                                                                                               Score 119.5; DB
Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABG20510 standard; protein; 100 AA.
Novel human diagnostic protein #20501.
WO200175067-A2.
                                                                                                                                                                                                                                                      protein; 207 AA. protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE55548 standard; protein; 203 AA.
Rat Protein Q63584, SEQ ID NO 1365.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADT60183 standard; protein; 244 AA.
Plant polypeptide, SEQ ID 10260.
US2004216190-A1.
28-OCT-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK66184 standard; protein; 119 AA
       10.2%;
27.3%;
                                                                                                                                                                                                               10.2%;
27.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                   10.1%;
20.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.1%;
20.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HAYA/) HAYASHIZAKI Y.
(KAMI/) KAMIYA M.
(SUZU/) SUZUKI T.
(HIRO/) HIROKAWA H.
                                                                                                                                                                                                                                                                       H. contortus pHC-2 pDE19704024-A1.16-APR-1998.(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse phX-6 protein.
US2004019006-A1.
   Query Match
Best Local Similarity
RESULT 922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
RESULT 928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
RESULT 929
                                                                                                                      Best Local Similarity RESULT 923
                                                                                                                                                                                                                       Best Local Similarity RESULT 924
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RESULT 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 927
                                                                                                                                                                                                                                                           AAW59051 standard;
                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NAKA/) NAKAO K.
(TANA/) TANAKA T.
                                                                                                                                                                                                06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2000.
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                                                                                                          Query Match
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RESULT 925
                                                                                                                                                                                                                                                           2222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human heat mitochondrial protein as a therapeutic target SeqID719, WO2003087768-A2.
    Score 123; DB 7; Length 219;
Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 123; DB 10; Length 219;
Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 13-MAY-2004.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 24.6%; Pred. No. 0.00049;
RESULT 917
                                                                                                                                    10.5%; Score 123; DB 7; Length 219; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                     Score 123; DB 7; Length 219; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                         Length 219;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAG16693 standard; protein; 204 AA.
Arabidopsis thallana protein fragment SEQ ID NO: 17437.
EP1033405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP52942 standard; protein; 218 AA.
Human lung specific protein sequence SEQ ID NO:185.
WO200264788-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEK18280 standard; protein; 219 AA.
Human transmembrane trafficking protein, TMP21.b.
US2006204503-A1.
                                                                                                                                                                                                                                                                                                                                                                                       Score 123; DB 8;
Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 123; DB 8;
Pred. No. 0.00049;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10.3%; Score 121; DB 5; 24.6%; Pred. No. 0.00079;
                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO protein; 219 AA. WO2004039956-A2.
                                             ADD48113 standard; protein; 219 AA.
Human Protein P49755, SEQ ID NO 14011.
WO2003016475-A2.
                                                                                                                                                                                 protein; 219 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP23066 standard; protein; 219 AA.
PRO polypeptide SEQ ID NO:160.
WO2004041170-A2.
                                                                                                                                                                                                                                                                                                               ADN03686 standard; protein; 219 AA.
                                                                                                                                                                                                                                                                                                                             Antipsoriatic protein sequence #40. WO2004028479-A2.
                                                                                                                                                                                                                  (MITO-) MITOKOR.

(BUCK-) BUCK INST AGE RES.

10.5%; Sc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PD 21-MAY-2004.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.6%;
RESULT 918
    10.5%;
24.6%;
                                                                                                                                                                                                                                                                                                                                                   PD 08-APR-2004.
PA (GETH ) GENENTECH INC.
Query Match 10.5%;
Best Local Similarity 24.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10.5%;
24.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-SEP-2006.
(BIOJ ) BIOGEN IDEC MA INC.
                                                                                     27-FEB-2003.
(GEHO ) GEN HOSPITAL CORP.
(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-AUG-2002.
(DIAD-) DIADEXUS INC.
Query Match
Best Local Similarity
RESULT 913
                                                                                                                                    Query Match
Best Local Similarity
RESULT 914
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RESULT 920
                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 915
                                                                                                                                                                                 ADJ68913 standard;
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(TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                EP1033405-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-SEP-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-FEB-200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 206;
                                                                                                                                                                                                                           82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 9.1%; Score 107; DB 8; Length 204; Local Similarity 19.8%; Pred. No. 0.021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8.8%; Score 103; DB 4; Length 427; 25.1%; Pred. No. 0.16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 237;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 211;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 214;
                     Length 244;
                                                                                                                                                    Length 204;
                                                                                                                                                                                                                                                                                                    Length 136;
                                                                                                                                                                                                    AEB88231 standard; protein; 136 AA.
Human protein highly expressed in cancer tissue SEQ ID NO
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG13953 standard; protein; 237 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 13636.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAG13954 standard; protein; 214 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 13637
EP1033405-A2.
06-SEP-2000.
                                                                         AAG07345 standard; protein; 204 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4463.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABB64424 standard; protein; 427 AA.
Drosophila melanogaster polypeptide SEQ ID NO 20064
                                                                                                                                                                                                                                                                                                                                                      ABP52926 standard; protein; 211 AA.
Human lung specific protein sequence SEQ ID NO:169.
WO200264788-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ63854 standard, protein, 204 AA.
Plant lipid metabolism protein Jb027 SEQ ID NO:56.
WO2004013304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADX95475 standard, protein; 102 AA.
Plant full length insert polypeptide segid 58139
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8.8%; Score 103; DB 3;
19.7%; Pred. No. 0.07;
                   9.3%; Score 110; DB 8; 27.7%; Pred. No. 0.013;
                                                                                                                                                                                                                                                        18-AUG-2005.
(RIVE-) FIVE PRIME THERAPEUTICS INC.
ery Match
9.2%; Score 108; DB 9;
ery Match
7.2%; Pred. No. 0.0093;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  9.2%; Score 108; DB 5; 23.8%; Pred. No. 0.018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8.8%; Score 103; DB 3;
19.7%; Pred. No. 0.06;
                                                                                                                                                  Score 109; DB 3;
Pred. No. 0.013;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AMPA-) FUNDACAO AMPARO A PESQUISA DO ESTADO.
ry Match
t Local Similarity 22.1%; Pred. No. 0.024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADY64838 standard; protein; 206 AA. S. mansoni protein SEQ ID 256. WO2005023979-A2.
                                                                                                                                                  9.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                   (DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
   (KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 933
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                                    Best Local Similarity
                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-SEP-2000.
                                                                                                                                06-SEP-2000.
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Best Local S
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                     Query Match
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ID ADX954
DE Plant
PN US2004
PD 19-FEE
PA (LIUJ/
PA (KOVA/
PA (SCRE/
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                                                       RESULT 931
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Length 102;
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(GEHU-) GERMAN HUMAN GENOME PROJECT.
ry Match
F Incal Similarity 24.2%; Pred. No. 1.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 8.2%; Score 96.5; DB 3; Length 621; Local Similarity 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 8.2%; Score 96.5; DB 4; Length 621;
Local Similarity 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 195;
                                                                                                                                            Score 101; DB 3; Length 214; Pred. No. 0.097;
                                                                                                                                                                                                                                                                                              Score 101; DB 3; Length 237; Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                               8.3%; Score 98; DB 3; Length 208; 21.2%; Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8:3%; Score 98; DB 3; Length 208; 21.2%; Pred. No. 0.19;
                                                                                                                                                                                                           AAG51637 standard; protein; 237 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 65557
EP1033405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                         AAGS4218 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 69107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABB68400 standard; protein; 195 AA.
Drosophila melanogaster polypeptide SEQ ID NO 31992.
WO200171042-A2.
Score 101.5; DB 8;
Pred. No. 0.03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAGO4572 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO:
EP1033405-A2.
                                                        A451638 standard, protein, 214 AA.
Arabidopsis thaliana protein fragment SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU53358 standard; protein; 544 AA.
Human testes-derived protein from DKFzphtes3_8g5
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU53359 standard; protein; 537 AA.
Human testes-derived DKFZphtes3_895 homologue.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8.2%; Score 96.5; DB 21.8%; Pred. No. 0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHU-) GERMAN HUMAN GENOME PROJECT.
ry Match
L Local Similarity 24.2%; Pred. No. 1.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human protein sequence SEQ ID NO:12237.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY83077 standard; protein; 621 AA.
F-box protein FBP-9.
WO200012679-A1.
    8.6%;
                                                                                                                                              Best Local Similarity 19.7%;
RESULT 941
                                                                                                                                                                                                                                                                                                   8.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAR-2000.
(UYNY ) UNIV NEW YORK STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-FEB-2001.
(HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 942
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 943
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  Query Match
Best Local Similarity
RESULT 940
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Human translocation associated protein designated Gp25L-H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 621;
                                                                                                      DB 5; Length 621;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8.1%; Score 95.5; DB 3; Length 169; 21.0%; Pred. No. 0.26;
                                                                                                                                                                                                                                                               Length 621;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 634;
                                                                                                                                                                                                                                                                                                                                                                                                                      8.2%; Score 96.5; DB 9; Length 621; 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 9; Length 628
                                                                                                                                                                                                                                                                                                                  ADX05819 standard; protein; 621 AA.
Cyclin-dependent kinase modulation biomarker SEQ ID NO 384.
WO2005012875-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADXO5821 standard; protein; 628 AA.
Cyclin-dependent kinase modulation biomarker SEQ ID NO 386.
WO2005012875-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAGI3955 standard; protein; 169 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 13638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8.2%; Score 96.5; DB 9; 24.2%; Pred. No. 1.4;
                                                                                                                                                                                                                                                             Score 96.5; DB 7;
Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                         ADB79933 standard; protein; 621 AA.
Human putative F-box protein Fbx21, SEQ ID 173
EP1279744-A2.
                                    PD 18-JUL-2002.
PA (UYNY ) UNIV NEW YORK STATE.
Query Match
Best Local Similarity 24.2%; Pred. No. 1.3;
RESULT 950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PA (BRIM') BRISTOL-MYERS SQUIBB CO.
Query Match
Best Local Similarity 24.2%; Pred. No. 1.4;
RESULT 955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8.2%; Score 96.5; Di 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.2%; Score 96.5; Di 24.2%; Pred. No. 1.4;
AAO22463 standard; protein; 621 AA.
Human F-box protein FBP9 SEQ ID No 28.
WO200255665-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AEA23925 standard; protein; 628 AA. Human PRO polypeptide SEQ ID NO 467, WO2005051988-A2. G9-JUN-2005. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEL92451 standard; protein; 621 AA. Human F-box protein FBP9. WO2006113251-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADY62385 standard; protein; 621 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU31647 standard, protein, 634 AA.
Novel human secreted protein #2138
WO200179449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW80293 standard; protein; 218 AA
                                                                                                                                                                                                                                                                                                                                                                               10-FEB-2005.
(BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                    PD 29-JAN-2003.
PA (WARN) WARNER LAMBERT CO.
Query Match
Best Local Similarity 24.2%;
RESULT 951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYNY ) UNIV NEW YORK STATE (CELG-) CELGENE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human F-box protein, FBP9.
WO2005020922-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 952
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RESULT 953
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Best Local Similarity
RESULT 954
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(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-2006
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Length 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 107;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 219;
                                                                                                                                         Length 218;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 93; DB 3; Length 493;
Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 93; DB 3; Length 493;
Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 493;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAG51639 standard; protein; 169 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 65559
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAG49571 standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 62725.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 лмыгызы standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 33594.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAG21783 standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 24466.
BP1033405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disease treating protein complex-derived protein #374 EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 7.9%; Score 93.5; DB 3;
Local Similarity 21.0%; Pred. No. 0.43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 7.9%; Score 93.5; DB 6; Best Local Similarity 19.8%; Pred. No. 0.62; RESULT 962
                                                                                                                                                                                                                                                                                                                                                                                                                           7.9%; Score 93.5; DB 5;
23.9%; Pred. No. 0.22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 93.5; DB 7;
Pred. No. 0.62;
Human ...
US5831052-A.
03-80V-1998.
(INCY-) INCYTE PHARM INC.
8.0%; Score 94; DB 2; ...
Match ...
13-71ty 21.4%; Pred. No. 0.54;
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Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 5;
                                                                                                                                                                                                                                         ABP61452 standard; protein; 107 AA.
Human NF-kB activating protein SEQ ID NO 57.
WO200253737-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.8%; Score 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat NOVX protein; 400 AA.
Rat NOVX protein homologue SegID 451
06.5PB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein sequence #SEQ ID 709.
20-Now Constitution  
20-Now Constit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK62496 standard; protein; 219 AA
                                                                                                                                                                                                                                                                                                                                         11-JUL-2002.
(ASAH ) ASAHI KASEI KOGYO KK.
Query Match 7.9%;
Best Local Similarity 23.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                           Best Local Similarity
RESULT 959
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(CELL-) CELLZOME AG.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 965
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-FEB-2002.
(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2000.
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                                                                                                                                         Query Match
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DB 9; Length 276;
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                                                                                                                                                                                                                  7.6%; Score 89; DB 3; Length 168; 24.3%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 89; DB 8; Length 223;
Pred. No. 1.9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 408;
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13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABG93808 standard; protein; 605 AA.
Transcription factor stress-related protein (TFSRP)
US2002102695-A1.
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Hyperthermophile Methanopyrus kandleri protein #679.
WO2003076575-A2.
                                                                                                                                                                                                                                                                                 ADY23730 standard; protein; 223 AA.
Plant full length insert polypeptide segid 71514.
US2004034888-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 7.6%; Score 89; DB 4 Local Similarity 37.7%; Pred. No. 4.5;
                                                                7.7%; Score 90.5; D. 24.7%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 87.5; D
Pred. No. 5.9;
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel human diagnostic protein; 408 AA. WO200175067-A2.
                                                                                                                       AAB27560 standard; protein; 168 AA.
Human secreted protein SEQ ID NO: 61.
WO200055175-A1.
C1-SEP-2000.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW47273 standard; protein; 1946 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU71799 standard; protein; 605 AA.
Zinc finger-3 (ZF-3).
WO200177311-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH78770 standard; protein; 384 AA
Pea GPA1 ortholog, SEQ ID 50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (FIDE-) FIDELITY SYSTEMS INC. (MALY/) MALYKH A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.4%;
Local Similarity 23.0%;
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Best Local Similarity 28.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SILVA) SILVA O D C E. (BOHN/) BOHNERT H J. (THIE/) THIELEN N V. (CHEN/) CHEN R.
                                                                                                                                                                                                                                                                                                                                                              (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                        (TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                                                                                                                      Best Local Similarity RESULT 978
                                                                                                                                                                                                                                                                                                                                                                                                                      SCREEN S E.
                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                          16-DEC-2004.
(RIKE ) RIKEN KK.
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        WO2004108930-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                  Query Match
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Mouse small endoplasmic reticulum resident protein 3 (smERp3) version #2.
EP1447412-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADV67442 standard; protein; 276 AA.
Amino acid sequence of a protein involved in FGF signal transduction.
                                                                                                                                               DB 6; Length 572;
                                                                                                                                                                                                                                                                                                       DB 7; Length 572;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 572;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.7%; Score 91; DB 5; Length 1450; 21.3%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.7%; Score 91; DB 8; Length 372; 24.2%; Pred. No. 2.4;
                                                                                                                                                                                                    ADK63196 standard; protein; 572 AA.
Disease treating protein complex-derived protein #724.
EP1338608-42.
27-AUG-2003.
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A. gossypii transport protein Oligo 167 SEQ ID 45.
WO2003014353-A2.
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7.7%; Score 90.5; DB 8;
Similarity 24.7%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH78766 standard; protein; 372 AA.
Nicotiana plumbaginifolia GPA1 ortholog, SEQ ID
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.7%; Score 90.5; Di 21.4%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.7%; Score 90.5; Di 24.7%; Pred. No. 1.8;
        23.0%; Pred. No. 2.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU02498 standard, protein, 276 AA.
Murine trinucleotide repeat protein (TRP).
                                                                                                                                                                                                                                                                                                     7.8%; Score 91.5; 22.0%; Pred. No. 4;
                                                                                                                                           7.8%; Score 91.5; 22.0%; Pred. No. 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 7.8%; Score 91.5; Best Local Similarity 22.0%; Pred. No. 4; RESULT 971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (INRG) INRA INST NAT RECH AGRONOMIQUE.
Query Match 7.7%; Score 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB55397 standard; protein; 1450 AA.
Lactococcus lactis protein yvcC.
                                                                                                                                                                                                                                                                                                                                                              ADS43610 standard; protein; 572 AA.
Bacterial polypeptide #22040.
US2003233675-A1.
                                            ABR53173 standard; protein; 572 AA Protein sequence #SEQ ID 1211.
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                                                                                                                                                                                                                                                                                                                                                                                                                      18-DEC-2003.
(CAOY/) CAO Y.
(HINK/) HINKLE G J.
(SLAT/) SLATER S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHEN/) CHEN X. (GOLD/) GOLDMAN B S.
                                                                                                                                                                                                                                                                                   (CELL-) CELLZOME AG.
                                                                                                                           (CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
RESULT 975
Best Local Similarity
RESULT 968
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RESULT 969
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Best Local Similarity
RESULT 976
                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BADI ) BASF AG.
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                                                                                    EP1258494-A1.
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                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                           RESULT 970
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Length 384;

., 8

#3

Length 605;

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AEA59839 standard; protein; 690 AA.
Streptococcus pneumoniae ORF amino acid sequence SEQ ID NO:4604
US2005136404-A1.
                                                                        DB 2; Length 1946;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABU02615 standard; protein; 690 AA.
S. pneumoniae type 4 strain protein from coding region #2193 W3200277021-A2.
                                                                                                                                                                                                               DB 8; Length 391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.3%; Score 86; DB 6; Length 491; 24.5%; Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                    DB 4; Length 488;
12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 86; DB 6; Length 690;
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7.3%; Score 86; DB 8; Length 690; 22.3%; Pred. No. 20;
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20;
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Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 86; DB 8; Length 687
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                ABU22666 standard; protein; 491 AA.
Protein encoded by Prokaryotic essential gene #8193.
WO200277183-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADR95969 standard; protein; 690 AA.
Novel S. pneumoniae protein sequence, SEQ ID 4604
US6800744-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM92253 standard; protein; 687 AA.
S pneumoniae antigenic protein sequence SeqID450.
W02004020609-A2.
(TUFT ) UNIV TUFTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptococcus pneumoniae protein, Seq ID No 4221. US6699703-B1.
                        PD 03-DEC-1997.
PA (NEST) SOC PROD NESTLE SA.
Query Match
Best Local Similarity 22.5%; Pred. No. 79;
RESULT 985
                                                                                                                                                                                                           Query Match 7.3%; Score 86;
Best Local Similarity 22.1%; Pred. No. RESULT 986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-0CT-2004.
(GENO-) GENOME THERAPEUTICS CORP.
(ery Match
7.3%; Score 86;
                                                                                                                                                                                                                                                                                                                                                    Score 86;
Pred. No.
Lactobacillus bulgaricus protease P.
                                                                                                                                                                                                                                                  Novel bone marrow polypeptide #46. 09-ARG-10151187-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK47706 standard; protein; 690 AA.
                                                                                                                    AD065925 standard; protein; 391 AA.
Novel human protein sequence #898.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAR-2004.
(GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                           28-JUL-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                  7.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIR-) CHIRON SPA.
(GENO-) INST GENOMIC RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-OCT-2002.
(ELIT-) ELITRA PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 993
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Best Local Similarity
RESULT 991
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Best Local Similarity
RESULT 992
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RESULT 989
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                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
RESULT 987
                                                                                                                                                                                                                                                                                                                09-AUG-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Length 690;
                                                                                                                                                                                                                                                                                                                                                                                      Length 690;
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Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 86; DB 10; Length 690;
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7.3%; Score 86; DB 6; Length 2301;
19.3%; Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 2938;
                                                                                                                                                                                                                                          Length 690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 690
AEJ71634 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7074914-B1.
                                                                                                                                                                                                                                                                                         AEJ85969 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7098023-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEL15538 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7122368-B1.
                                                                                                                                            ABJ78609 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221
US7081530-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                            ALLUGZBB standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.057115731-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEL53946 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221
                                                                                           DB 10;
20;
                                                                                                                                                                                                                                        DB 10;
20;
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20;
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Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                   Score 86; DB 10;
Pred. No. 20;
                                           PD 11-JUL-2006.
PA (SNFI) SANOFI PASTEUR LTEE.
Query Match 7.3%; Score 86;
Best Local Similarity 22.3%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 86;
Pred. No.
                                                                                                                                                                                                                                          Score 86;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABUS2379 standard; protein; 2301 AA.
Human GPCR related protein NOV29a.
WO200279398-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR59923 standard; protein; 2938 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADN18817 standard; protein; 3092 AA
                                                                                                                                                                                                                                                                                                                                           29-70-2006.
(SNFI ) SANOFI PASTEUR LTD.
7.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 22.3%;
RESULT 998
                                                                                                                                                                                                    25-JUL-2006.

(SMRI) SANOFI PASTEUR LIEE.

Query, Match

Best Local Similarity 22.3%;
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7.3%;
tr Local Similarity 22.3%;
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Local Similarity 22.3%;
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Best Local Similarity 19.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacterial polypeptide #1470.
US2003233675-Al.
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(SNFI ) SANOFI PASTEUR LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2006.
(SNFI ) SANOFI PASTEUR LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SCHE ) SCHERING CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
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HINKLE G J.
SLATER S C.
CHEN X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GOLDMAN B S
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WO9416069-A2.
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(CHEN/) C
(GOLD/) C
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RESULT 995
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22-FEB-2001

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ABUS1080 standard; protein; 711 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #13.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                     AABS6790 standard; protein; 173 AA.
Human prostate cancer antigen protein sequence SEQ ID NO:1368.
WO200055174-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 384;
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                                                                                                                                                                                                                                                                                                                DB 3; Length 147;
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                                                                                                                                                       7.3%; Score 85.5; DB 3; Length 147; 25.2%; Pred. No. 2.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 173;
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Length 3092;
                                                                                                                                                                                                             AAB27611 standard; protein; 147 AA.
Human secreted protein BLAST search protein SEQ ID NO: 112.
WO200055175-A1.
                                                        AAB27610 standard; protein; 147 AA.
Human secreted protein BLAST search protein SEQ ID NO: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG21784 standard; protein; 384 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 24467.
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAG49572 standard; protein; 384 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 62726.
EP1031405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAG28394 standard; protein; 384 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 33595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.3%; Score 85.5; DB 3;
19.4%; Pred. No. 9.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.3%; Score 85.5; DB 3; 25.2%; Pred. No. 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.3%; Score 85.5; DB 3; 19.4%; Pred. No. 9.6;
7.3%; Score 86; DB 8; I
33.7%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADW07163 standard; protein; 601 AA.
Bos taurus keratin 5 protein sequence SeqID3
NZS25023-A.
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19.4%; Pred. No. 9.6;
                                                                                                                                                                                                                                                                                                              7.3%; Score 85.5; D 25.2%; Pred. No. 2.4;
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18.2%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB77434 standard; protein; 623 AA. Human tumour marker protein se20-7.WO200238803-A2.
                                                                                            WO200055175-A1.
21-SEP-2000.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                        21-SEP-2000.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (STAN) STHERLAND G T.
(STAN) STANFIELD A M.
(SPEL) SPELMAN R J.
(ANKE) ANKERSMIT-UDY A E L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC. (ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 1
RESULT 1010
ID ABUSJO80 standard, pro
DE Intracellular traffick
PN WO200112659-A2.
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                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 1004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
              Best Local Similarity RESULT 1002
                                                                                                                                                                     Local Similarity
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  Query Match
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ABUS3072 standard; protein; 941 AA.
Intracellular trafficking-associated DKFZphtes3_1913 homologue #5.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                    ABU53070 standard; protein; 951 AA.
Intracellular trafficking-associated DKFZphtes3_1913 homologue #3.
                                                                       JULIAN Standard; protein; 907 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #6.
WO200112659-A2.
22-FEB-2001.
(GEHU-) GERMAN HUMAN GENOME PROJECT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABUS3071 standard; protein; 953 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #4.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABU53069 standard; protein; 953 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #2.
WO200112659-A2.
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                                                                                                                                                                               Score 85.5; DB 4; Length 907;
Pred. No. 33;
                Length 711;
                                                                                                                                                                                                                                                                                                                                              Length 941;
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Protein differentially regulated in prostate cancer #48
                  DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 7.3%; Score 85.5; DB 6; Best Local Similarity 18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4;
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Pred. No. 1.2e+02;
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18.2%; Pred. No. 1.2e+02;
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Human golgin-245 splice variant 2 polypeptide.
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR61599 standard; protein; 2228 AA.
Human golgin-245 splice variant 1 polypeptide.
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR61601 standard; protein; 2250 AA.
Human golgin-245 splice variant 3 polypeptide.
WO2003087403-A2.
SOCT-2003.
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          Best Local Similarity 18.2%; Fred. No. 23; RESULT 1011
                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 18.2%; Pred. No. 35; RESULT 1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 85.5; Pred. No. 36;
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Pred. No. 36;
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Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHU-) GERMAN HUMAN GENOME PROJECT.
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(GEHU-) GERMAN HUMAN GENOME PROJECT.
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GERMAN HUMAN GENOME PROJECT
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(EVOT-) EVOTEC NEUROSCIENCES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EVOT-) EVOTEC NEUROSCIENCES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ORIG-) ORIGENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 7.3%;
Best Local Similarity 18.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 18.2%;
RESULT 1017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 7.3%;
Best Local Similarity 18.2%;
RESULT 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 7.3%;
Best Local Similarity 18.2%;
                                                                                                                                                                                 Query Match 7.3%;
Best Local Similarity 18.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200281638-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-OCT-2003
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02-MAY-2002.
(CHIR-) CHIRON SPA.
(GENO-) INST GENOMIC RES.
                                         (SLAT/) SLATER S C. (CHEN/) CHEN X. (GOLD/) GOLDMAN B S.
                      HINKLE G J.
SLATER S C.
                                                                                         Query Match
Best Local Similarity
RESULT 1028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-SEP-2001
(PEKE ) PE CORP NY.
                                                                                                                                                   ADB12159 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABB70375 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-2004.
(KEKU/) KEKUDA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FR2824074-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                     Match 7.3%; Score 85.5; DB 8; Length 4447; Local Similarity 24.4%; Pred. No. 3.3e+02;
                                                                                                                                                                  Length 2252;
                        Length 2250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.2%; Score 84.5; DB 7; Length 773; 21.1%; Pred. No. 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.2%; Score 85; DB 8; Length 223; 22.0%; Pred. No. 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4; Length 419;
12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 722,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADK63862 standard; protein; 773 AA.
Disease treating protein complex-derived protein #1444.
EP1338608-A2.
                                                                                                          PD 23-OCT-2003.
PA (EVOT-) EVOTEC NEUROSCIENCES GMBH.
Query Match 7.3%; Score 85.5; DB 7;
Best Local Similarity 18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ADY10508 standard, protein; 223 AA.
Plant full length insert polypeptide segid 66323.
US2004034888-A1.
Query Match 7.3%; Score 85.5; DB 7; Best Local Similarity 18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.2%; Score 84.5; DB 6; 21.1%; Pred. No. 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 03-APR-2003.

PA (INCY-) INCYTE GENOMICS INC.

Query Match 7.2%; Score 85; DB 6;

Best Local Similarity 32.2%; Pred. No. 27;

RESULT 1025
                                                                     ABR61602 standard; protein; 2252 AA.
Human golgin-245 splice variant 4 polypeptide.
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.2%; Score 85; 22.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG13105 standard; protein; 419 AA.
Novel human diagnostic protein #13096.
WO200175067-A2.
                                                                                                                                                                                                       Bacterial polypeptide #23143.
18-DECT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 722 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR53767 standard; protein; 773 AA.
Protein sequence #SEQ ID 2399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADN18818 standard; protein; 773 AA.
Bacterial polypeptide #1471.
US2003233675-A1.
18-DEC-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lu-) CELLZOME AG.
Lu-Y Match
Best Local Similarity 2.
RESULT 1027
ID ADN18918 stand-
DE Bacterial
PN US2nnnnn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR69620 standard; pro
Human CGDD-20 protein.
WO2003027263-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 1026
                                                                                                                                                                                                                                                                                                                                                                (GOLD/) GOLDMAN B S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 1024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 1023
                                                                                                                                                                                                                                                                                         (CAOY/) CAO Y.
(HINK/) HINKLE G J.
(SLAT/) SLATER S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-OCT-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                   (CHEN/) CHEN X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EP1258494-A1.
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Si
RESULT 1022
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DB 8; Length 773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 84; DB 4; Length 1605; Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 486;
                                                                                                                                                       Length 227;
                                                                                                                                                                                                                                                                                                                                Match 7.1%; Score 84; DB 5; Length 486; Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'Match 7.1%; Score 84; DB 8; Length 486; Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 486;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 827;
                                                                                                                                                                                                                                                                                                                                                                                         ADV88989 standard; protein; 486 AA.
Streptococcus agalactiae protein sequence, SEQ ID 1383
                                                                        Alloiococcus otitis antigenic protein SEQ ID NO:6554 WO2003048304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Drosophila melanogaëter polypeptide SEQ ID NO 37917.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC31334 standard; protein; 827 AA.
Human novel polypeptide sequence, SEQ ID NO:1416.
WO2003029271-A2.
                                                                                                                                                         9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
8
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADV80242 standard; protein; 486 AA.
Streptococcus agalactiae protein, SEQ ID 1383
WO200292818-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptococcus agalactiae protein, 486 AA.
WO200292818-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB -
                                                                                                                                                       Score 84; DB (
Pred. No. 6.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 7.1%; Score 84; DB Best Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7.1%; Score 84; DB 20.5%; Pred. No. 19;
Score 84.5; 1
Pred. No. 34;
                                                                                                                                                                                                           ABP27911 standard; protein; 486 AA.
Streptococcus polypeptide SEQ ID NO 4798.
WO200234711-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.1%; Score 84; 32.2%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL24093 standard; protein; 2301 AA.
Human NOVX polypeptide #69.
US2004002120-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; 1605 AA
                                                        protein; 227 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT RECH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-NOV-2002.
(INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT TECH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-NOV-2002.
(INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT FECH SCI.
7.2%;
                                                                                                                                                     Query Match 7.1%;
Best Local Similarity 19.6%;
RESULT 1029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.1%;
23.6%;
                                                                                                                12-JUN-2003.
(AMHP ) WYETH HOLDINGS CORP.
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Human protein encoded by clone THYMU20098350.
EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UNIV NORTH CAROLINA STATE.

7.1%; Score 83.5; DB 9; Length 406; Similarity 18.2%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%; Score 83.5; DB 4; Length 399; 19.7%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 8; Length 326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 8; Length 2301;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB75224 standard; protein; 399 AA.
Drosophila gustatory receptor GR59E.2 protein sequence.
WO200077208-A2.
21-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADX96054 standard; protein; 326 AA.
Plant full length insert polypeptide segid 58718.
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 7.1%; Score 84; DB 8; 1
Best Local Similarity 18.3%; Pred. No. 1.8e+02;
RESULT 1036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAE30522 standard; protein; 399 AA.
Fruit fly gustatory receptor protein, Gr59E2.
WO200268593-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%; Score 83.5; 1
21.3%; Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.1%; Score 83.5; 1
19.7%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEC57546 standard; protein; 406 AA.
L. acidophilus ABC transporter permease.
WO2005084411-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB65618 standard; protein; 567 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                GORMAN L.
MALYANKAR U M.
BOLDOG F L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SHIMKETS R A.
RASTELLI L.
SPADERNA S K.
LAROCHELLE W J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KHRAMTSOV N V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-FEB-2004.
(LIUJ/) LIU J.
(ZHOJ/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                 POCHART P F. FERNANDES E R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HERRMANN J L.
     TCHERNEV V T
                                                                                               MERNET C A M
                                                                                                                                                                                                        SHENOY S G.
PADIGARU M.
TAUPIER R J.
MILLER C E.
CASMAN S J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 1038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
RESULT 1040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 1039
                                                                                                                                                                                                                                                                                                                                                           SMITHSON G
                                                                                                                                                                                                                                                                                                    PENA C E A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                                                                                                          GUSEV V Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                   (MALY/)
(BOLD/)
(GUOX/
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AEB56515 standard; protein; 1022 AA.
Radiochemotherapy response detection associated protein SEQ ID NO 90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-AUG-2005.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                         Length 567;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 8; Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 7.1%; Score 83.5; DB 4; Length 914; Local 'Similarity 19.7%; Pred. No. 55;
                                                                                                                                                                                                                                                                     Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 8; Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ37107 standard; protein; 590 AA.

Human malignant pleural mesothelioma (MPM) protein #6.
US2003129760-A1.

(BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB68295 standard; protein; 914 AA.
Drosophila melanogaster polypeptide SEQ ID NO 31677.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 9;
                                                                                                                                                                                                                                                                DB 7;
                                                                         DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 7.1%; Score 83.5; DB 9;
Best Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (IPSO-) IPSOGEN.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
(PAOL-) INST PAOLI CALMETTES.
                                                                       Query Match 7.1%; Score 83.5; 1
Best Local Similarity 22.5%; Pred. No. 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                  (MILL-) MILLENNIUM PHARM INC.
ry Match 7.1%; Score 83.5; I
t Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 7.1%; Score 83.5; I
Best Local Similarity 25.1%; Pred. No. 64;
                                                                                                                                                                                                                                                                Query Match 7.1%; Score 83.5; 1
Best Local Similarity 22.5%; Pred. No. 29;
RESULT 1042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 7.1%; Score 83.5; I Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 7.1%; Score 83.5;
Best Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEB87738 standard; protein; 590 AA.
Human cytokeratin 5, breast tumor marker.
WO2005071419-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .1%; Score 83.5;
                                                                                                                                                                                                                   13-MAR-2003.
(BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1044
ID ADR99136 standard; protein; 590 AA.
DE Keratin 5, KRT5, SEQ ID 142.
PN WO2004078035-A2.
                                                                                                                                                                                                                                                                                                                                     ADB75394 standard; protein; 590 AA.
Prostate cancer marker protein.
WO2003009814-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEA15495 standard, protein, 590 AA.
Human polypeptide #114.
WO2005047534-A2.
                                                                                                                                            protein; 590 AA
07-MAY-2003.
(HELL-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 22.5%;
RESULT 1045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FARB ) BAYER HEALTHCARE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FARB ) BAYER PHARM CORP
                                                                                                                                                                    Keratin 5 SEQ ID NO:20.
WO2003021229-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                            ADB70328 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                 06-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2005
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14-APR-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-200
                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADR49442 standard; protein; 276 AA.
Mouse small endoplasmic reticulum resident protein 3 (smERp3) version #1.
EP1447412-A1.
                                                                                                                                                      AELS7159 standard; protein; 1062 AA.
Human SLIT-ROBO Rho GTPase activating protein 1, SEQ ID NO: 1622.
US2006216722-A1.
28-SEP-2006.
ADA13347 standard; protein; 1062 AA.
Human intracellular signalling molecule INTSIG-29, SEQ ID NO:29.
WO2003031568-A2.
                                                                                                                                                                                                                                                                                                                                    7.1%; Score 83.5; DB 10; Length 1062; 25.1%; Pred. No. 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AED11485 standard; protein; 181 AA.
Human gene 16 encoded secreted protein HIGAN47, SEQ ID NO: 46.
US2005214786-A1.
                                                                                                    DB 6; Length 1062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ92059 standard; protein; 474 AA.
Soybean hydroperoxide lyase from clone sdp3c.pk017.j17.fis
US2004010822-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 7; Length 181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7.1%; Score 83; DB 8; Length 282; 24.4%; Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 83; DB 8; Length 474;
Pred. No. 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.1%; Score 83; DB 8; Length 276; 25.0%; Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 83; DB 8; Length 429;
Pred. No. 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADY23081 standard, protein; 282 AA.
Plant full length insert polypeptide seqid 70865.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 9;
                                                                                                7.1%; Score 83.5; 1
25.1%; Pred. No. 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                          MOGNOS OF SEP-2003.

(HUMA-) HUMAN GENOME SCI INC.

7.1%; Score 83; DB

7.1%; Pred. No. 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USEC-2005.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) Autch
7.1%; Score 83;
                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted protein SEQ ID NO:46 WO2003072761-A1.
                                                                                                                                                                                                                                                                                                                                                                                          ADC78239 standard; protein; 181 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADT58337 standard; protein; 429 AA Plant polypeptide, SEQ ID 8414. US2004216190-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-AUG-2004.
(UYUT-) UNIV UTRECHT HOLDING BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%;
24.5%;
                                                                              (INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.1%;
                                                                                                                                                                                                                                  (BETS/) BETSHOLTZ C.
(TRYG/) TRYGGVASON K.
(TAKE)/ TAKEMOTO M.
(HELL/) HE L.
(PATR/) PATRAKKAS J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LIUJ/) LIU J.
(ZOVA/) ZOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
SSULT 1052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-OCT-2004.
(KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-2004.
(MCGO/) MCGONIGLE B.
                                                                                              Query Match
Best Local Similarity
RESULT 1050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
RESULT 1053
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RESULT 1055
                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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RESULT 1054
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Best Local Similarity
RESULT 1056
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                                                            17-APR-2003
                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                      RESULT 1051
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DB 8; Length 325;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 9; Length 403;
                                                                                                               Length 474;
                                                                                                                                                                                                                                                                                                                                                                            Score 83; DB 8; Length 489;
Pred. No. 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.0%; Score 82; DB 7; Length 211; 23.6%; Pred. No. 9.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Length 211;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse squalene epoxidase (SPF) protein sequence SeqID2 JP2005095162-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK62500 standard; protein; 211 AA.
Disease treating protein complex-derived protein #376.
EP1338608-A2.
DE Soybean hydroperoxide lyase SEQ ID NO:11.

PN WO2005089198-A2.

PD 29-SEP-2005.

PA (DUPO) DU PONT DE NEMOURS & CO E I.

Query Match

RESULT 1058

ID ADA78603 standard; protein; 489 AA.

PN USZO004034888-A1.

PN USZO04034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB70282 standard; protein; 984 AA.
C. neoformans amino acid sequence SEQ ID NO:3326.
WO2003052076-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.0%; Score 82; DB (23.6%; Pred. No. 9.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP04221 standard; protein; 325 AA.
Human colon specific protein SEQ ID NO:202.
WO2004050858-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 82.5; Pred. No. 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 82.5; I
Pred. No. 21;
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Pred. No. 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADI16914 standard; protein; 403 AA.
Murine NOVX protein homologue SeqID 450.
WO200268649-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABRS2924 standard; protein; 211 AA.
Protein sequence #SEQ ID 713.
EP1258494-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 403 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH78758 standard; protein; 366 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 7.0%;
Best Local Similarity 22.6%;
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Best Local Similarity 22.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ARAI/) ARAI H.
(SHIB/) SHIBATA N.
(CHUS ) CHUGAI PHARM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                              24.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ELIT-) ELITRA PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUN-2004.
(DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                      (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                             (SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CELL-) CELLZOME AG.
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RESULT 1065
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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                      DB 6; Length 610; 44;
                                                                                                                                                                                                                                                                    DB 6; Length 610;
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                                                                                                                                                                                                                                                                                                                                                                                      Length 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6; Length 610;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Length 610
                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human secreted and transmembrane protein PRO872.
US203036144-Al.
                                                                                                           ABU89872 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872
20.003034147-A1.
                                                                                                                                                                                                           ABR68121 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027264-A1.
06-PEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR94650 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003044926-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR74888 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane protein (PRO) #46. US2003036149-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO08682 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044923-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein (PRO) #46. US2003040062-A1.
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                                                                                                                                                                                                                                                                                                                                                                                DB
44;
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44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
44;
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Human secreted/transmembrane protein, #41.
US2002160384-A1.
                                                                                                                                                                      Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.0%; Score 82; 26.3%; Pred. No.
             ABU82751 standard; protein; 610 AA.
Human PRO polypeptide #46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU92605 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU13882 standard; protein; 610 AA
                                                                                                                                                                                                                                                                      7.0%;
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                                                                                                                                                                      7.0%;
26.3%;
                                                                    7.0%;
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26.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.0%;
26.3%;
                                                                                                                                                                                                                                                                                                                                                       31-OCT-2002 (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 1086
                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 1087
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RESULT 1092
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 1089
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Best Local Similarity
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RESULT 1091
                                                                                 Best Local Similarity
                                        US2003032113-Ā1
13-FEB-2003.
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                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                              RESULT 1085
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RESULT 1084
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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                           DB 6; Length 610;
44;
                                                                                                                                                 Length 610;
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                                                                                                                                                                                                                                                                                                                        Length 610;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 610;
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                                                                                                                                                                                                                                                                                                                                         Novel human secreted and transmembrane protein PRO872. US2003027277-A1.
                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO872. US2003013153-A1.
                                                                                                                                                                                                                                                             мвиу/уув standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003017544-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human secreted and transmembrane protein PRO872 US2003003531-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane protein (PRO) #46. US2003036162-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted transmembrane protein (PRO) #46. US2003036146-Al.
                                                                                                                                                                                                                                                                                                                        DB 6;
44;
                                                                                                                                                 9
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9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6;
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6
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44;
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44;
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44;
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44;
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                                                                                                                                                                                                                                    Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                        Score 82;
Pred. No.
                                                                                                                                                 Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 82;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #46.
US2003036141-Al.
                                                                           Human PRO polypeptide #46.
20-FFB.
                                                                                                                                                                                     protein; 610 AA
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                                                                                                                                                7.0%;
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                                                Query Match 7.0%;
Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Human PRO872 polypeptide.
US2002103125-A1.
01-AUG-2002.
(GETH ) GENENTECH LTD.
                                                                                                                        20-FEB-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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US2003036137-A1.
                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                   ABU98783 standard;
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6; Length 610; 6; Length 610; DB 6; Length 610; 44; Score 82; DB 6; Length 610; Pred. No. 44; DB 6; Length 610; 44; Length 610; Length 610; Length 610; Length 610; Length 610, ABUB5008 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872. US2003032114-A1. AB000147 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872. US2003032101-A1. ABR98787 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040064-A1. 27-FEB-2003. ABR78272 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003054474-A1. ABR92210 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003036160-A1. LT 1104 ABR99397 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. AB002124 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003040054-A1. 27-FEB-2003. ABO16310 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003027267-A1. ABO18851 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003044925-A1. ABO11479 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003036124-A1. 20-FEB-2003. . 9 .. 9 . 9 . 9 DB 44; 26.3%; Pred. No. 44; Score 82; Pred. No. 4 Score 82; Pred. No. 4 Score 82; Pred. No. 7.0%; 7.0%; 7.0%; 7.0%; 7.0%; 26.3%; 7.0%; 7.0%; 7.0%; 26.3%; 7.0%; 26.3%; 7.0%; 26.3%; 20-MAR-2003. (GETH) GENENTECH INC. Query Match Best Local Similarity RESULT 1110 Query Match Best Local Similarity Best Local Similarity RESULT 1104 Query Match Best Local Similarity Query Match Best Local Similarity RESULT 1108 Best Local Similarity Best Local Similarity Best Local Similarity RESULT 1114 Best Local Similarity RESULT 1106 Best Local Similarity Best Local Similarity US2003040063-A1. 27-FEB-2003. 06-FEB-2003. 20-FEB-2003. 13-FEB-2003. 06-MAR-2003. 13-FEB-2003. Query Match Query Match Query Match Query Match Query Match Query Match RESULT 1105 ID ABR98787 DE Human se PN US200304 PD 27-FEB-2 RESULT 1109 RESULT 1111 RESULT 1107 RESULT 1112 RESULT PA DE PN PN

DB 6; Length 610; 44; DB 6; Length 610; 44; Length 610; Length 610; Length 610; Length 610; Length 610; Length 610; Length 610 Length 610; ABU88698 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872. US2003036133-A1. Novel human secreted and transmembrane protein PRO872. US2003022294-A1. AB019156 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872 US2003036118-A1. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003027275-Al. Human secreted polypeptide PRO872, SEQ ID NO:92.02003036148-Al. ABU65614 standard; protein; 610 AA. Human secreted/transmembrane protein, SEQ ID 92. Human secreted/transmembrane protein (PRO) #46. US2003027324-A1. Human secreted/transmembrane protein (PRO) #46. US2003036134-Al. ABO11174 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003036123-A1. 20-FBB-2003. ABO16005 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003040060-A1. ABO13711 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46 US2003044916-A1. DB 6; 44; DB 6; 9 9 9 DB 6; DB 6; 44; . 9 DB / 44; DB / DB 44; DB 44; 44; Score 82; Pred. No. 7.0%; Score 82; 26.3%; Pred. No. Score 82; Pred. No. Score 82; Pred. No. protein; 610 AA protein; 610 AA 7.0%; 7.0%; 7.0%; 7.0%; 26.3%; 7.0%; 26.3%; 7.0%; 7.0%; 26.3%; 7.0%; 7.0%; 20-FEB-2003. (GETH) GENENTECH INC. Best Local Similarity RESULT 1117 Best Local Similarity RESULT 1118 Best Local Similarity Best Local Similarity RESULT 1115 ABU83393 standard; Best Local Similarity RESULT 1116 ABO06194 standard; Best Local Similarity Best Local Similarity Best Local Similarity Local Similarity 20-FEB-2003. 20-FEB-2003. 30-JAN-2003. 27-FEB-2003. Ouery Match Query Match Ouery Match Query Match

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(GETH ) GENENTECH INC.
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                           Length 610;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Novel human secreted and transmembrane protein PRO872.
US2003036117-A1.
                                                                                                                                                                                                                                                                   ABR67097 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US200302266-A1.
06-F8E-2003.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032138-A1.
13-FEB-2003.
                                                                                                                           DB 6;
44;
                           Score 82; DB 6;
Pred. No. 44;
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44;
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                                                                                                                                                                  ÀBO03649 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003036128-A1.
20-FEB-2003.
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26.3%; Pred. No. 44;
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Human secreted/transmembrane protein, PRO872
US2003022298-A1.
                                                                                                                                                                                                                                                                                                                                                             ABO15700 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO)
US2003054483-A1.
                                                                                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
                                                            Human PRO polypeptide #46.
US2003032117-A1.
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US2003032130-Al.
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Human PRO polypeptide #46.
US2003032102-A1.
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Human PR0872 protein.
US2003036143-Al.
20-FEB-2003.
                             7.0%;
26.3%;
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26.3%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1127
                                     Best Local Similarity
RESULT 1125
                                                                                                                         Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1128
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Best Local Similarity
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Best Local Similarity
RESULT 1130
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RESULT 1133
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RESULT 1134
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US2003036156-A1 20-FEB-2003.
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                             Query Match
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                                                                                                                                                      RESULT 1126
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Score 82; DB 6; Length 610; Pred. No. 44;
 Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                        Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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44;
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44;
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44;
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                                          ABR69341 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036132-A1.
                                                                                                                                                                                                                                                                                                                                                                                     ABR60081 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032137-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR67816 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR65204 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027268-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR68426 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027274-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR71838 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032135-A1.
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44;
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44;
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Human secreted/transmembrane protein, #41.
US2003027162-A1.
                                                                                                                                                                                                                                 Score 82;
Pred. No.
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26.3%; Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                         protein; 610 AA.
                                                                                                                                                                                                                                                                             ABU81284 standard; protein; 610 AA.
Human PRO polypeptide #46.
US2003017542-A1.
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26.3%;
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26.3%;
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Best Local Similarity 26.3%;
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US2003008353-A1.
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(GETH ) GENENTECH INC.
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RESULT 1140
Query Match
Best Local Similarity
                                                                                                                       Local Similarity
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Best Local Similarity
RESULT 1138
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                                                                                                                                                       ABO01482 standard;
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RESULT 1142
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                                                                                                                                       RESULT 1136
                               RESULT 1135
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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                       7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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44;
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Pred. No. 44;
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Pred. No. 44;
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44;
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44;
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Novel human secreted and transmembrane protein PRO872.
US2003032119-A1.
                                                                                                                                                                                                                                                                                              ABU94944 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US200307263-Al.
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RESULT 1153
ID ABR68731 standard; protein; 610 AA.
DE Human secreted polypeptide PRO872, SEQ ID NO:92.
PN US2003027271-A1.
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US2003036125-A1.
                                                                                                                                                                                                                                                                                                                                                                  DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 26.3%; Pred. No. 44;
RESULT 1149
ID ABU84003 standard; protein; 610 AA.
BD Human secreted/transmembrane protein (PRO) #46.
PN US2003032111-A1.
PD 13-FEB-2003.
                                                                                                                                                                        ABU83088 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032105-A1.
                                                  ABU89008 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003022297-A1.
                                                                                                                                                                                                                                                                                                                                                                    7.0%; Score 82; 26.3%; Pred. No.
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Pred. No.
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Pred. No.
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6.3%; Pred. No.
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Human PRO872 polypeptide.
US2002127576-A1.
12-SEP-2002.
(GETH ) GENENTECH INC.
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26.3%;
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RESULT 1151
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RESULT 1150
ID ABU93654 standard; pr
DE Novel human secreted
PN US2003032119-A1.
PD 13-FEB-2003.
PA (GETH ) GENENTECH INC
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Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1154
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                                   RESULT 1145
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DB 6; Length 610;
44;
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Novel human secreted and transmembrane protein PRO872.
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Novel human secreted and transmembrane protein PRO872.
20-FBB-2003.
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Novel human secreted and transmembrane protein PRO872.
US2003032104-A1.
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                        ABR99092 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
USO03040068-A1.
27-FEB-2003.
                                                                        DB 6;
44;
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Pred. No.
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Pred. No.
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US2003027280-Al.
                                                                                                                                                                                                                                                                                                                                                                       protein; 610 AA
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Best Local Similarity 26.3%;
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Human PRO polypeptide #46.
US2003036138-Al.
20-FEB-2003.
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Human PRO polypeptide #46.
US2003040066-A1.
Best Local Similarity
RESULT 1155
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Best Local Similarity
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6; Length 610; Score 82; DB 6; Length 610; Pred. No. 44; Length 610; 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44; Length 610; Score 82; DB 6; Length 610; Pred. No. 44; Score 82; DB 6; Length 610; Pred. No. 44; DB 6; Length 610; 44; 6; Length 610; Score 82; DB 6; Length 610; Pred. No. 44; Novel human secreted and transmembrane protein PRO872 Novel human secreted and transmembrane protein PRO872 US2003022296-A1. คอบชอช48 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872. US_DD12131-A1. ABR90905 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040058-A1. Human secreted polypeptide PRO872, SEQ ID NO:92. Human secreted/transmembrane protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032129-A1. DB 6; DB 6; DB 6 44; DB 44; Human sectreted/transmembrane protein, #41. US2002142961-A1. 03-0CT-2002 (GETH) GENENTECH INC. Score 82; DB Pred. No. 44; Score 82; Pred. No. Score 82; Pred. No. 7.0%; Score 82; 26.3%; Pred. No. Score 82; Pred. No. Human PRO polypeptide #46. Human PRO polypeptide #46.
US2003032106-Al.
113.FEB-2003. protein; 610 AA. ABU94637 standard; protein; 610 AA. Human PRO polypeptide #46. US2003032103-A1. ABU58922 standard; protein; 610 AA ABO04564 standard; protein; 610 AA. Human PRO polypeptide #46. 7.0%; 26.3%; 7.0%; 7.0%; 7.0%; 7.0%; 26.3%; 7.0%; 26.3%; 7.0%; 7.0%; 06-FEB-2003. (GETH) GENENTECH INC. Query Match Best Local Similarity RESULT 1168 Best Local Similarity RESULT 1169 Query Match Best Local Similarity Best Local Similarity RESULT 1167 Query Match Best Local Similarity RESULT 1171 Query Match Best Local Similarity Best Local Similarity RESULT 1166 Ouery Match Best Local Similarity RESULT 1172 Query Match Best Local Similarity Best Local Similarity ABU99817 standard; US2003036155-A1. 20-FEB-2003. 30-JAN-2003. 27-FEB-2003. 23-JAN-2003. 13-FEB-2003. 13-FEB-2003. Query Match Query Match Query Match RESULT 1170 RESULT 1174 RESULT

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Novel human secreted and transmembrane protein PRO872.
US2003022187-A1.
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Novel human secreted and transmembrane protein PRO872.
US2003036154-A1.
20-PEB-2001
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Novel human secreted or transmembrane protein PRO1063 US2003027985-A1.
                                          Human secreted polypeptide PRO872, SEQ ID NO:92. US2003032139-Al. 13-PEB-2003.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003036165-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027262-A1.
                        DB 6;
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Human secreted/transmembrane protein (PRO) #46.
US2003036142-A1.
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44;
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Pred. No.
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Pred. No.
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US2003032110-Al.
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US2003022301-A1.
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US2003036145-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
                                    Best Local Similarity
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                                                                                                                                                                                                                                         ABU98478 standard;
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ID ABU95869 standard;
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US2003032107-A1.
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                        Query Match
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ABU93349 standard; protein; 610 AA.
Human PRO polypeptide #46.
US2003017541-A1.
 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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Pred. No. 44;
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44;
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Pred. No. 44;
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Novel human secreted and transmembrane protein PRO872.
23-03017476-A1.
                                     ABU90182 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036153-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                             ABR70923 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003040069-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003032122-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003032116-A1.
                                                                                                                                                AB003597 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044931-A1.
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Human secreted/transmembrane protein (PRO) #46
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44;
                                                                                                        7.0%; Score 82; DB 26.3%; Pred. No. 44;
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Pred. No. 4
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Human PRO protein #46.
US2003036139-A1.
20-FEB-2003.
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26.3%;
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Best Local Similarity
RESULT 1188
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RESULT 1195
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RESULT 1196
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Best Local Similarity
RESULT 1193
               Best Local Similarity
                                                                                                                    Best Local Similarity RESULT 1187
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Best Local Similarity
RESULT 1192
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                                                                                                                                                                                                                                                                                     US2003036150-A1.
20-FEB-2003.
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Length 610;
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            Length 610;
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Novel human secreted and transmembrane protein PRO872.
US2003008322-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU81589 standard, protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2002177164-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR70618 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040076-A1.
            DB 6;
44;
                                                            Human secreted/transmembrane protein (PRO) #46.
US2003017543-A1.
                                                                                                                                                Human secreted/transmembrane protein (PRO) #46 US2003036152-A1.
                                                                                                                                                                                          DB 6;
44;
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44;
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44;
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Human secreted/transmembrane protein #46.
US2002127584-A1.
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(GETH ) GENENTECH INC.
7.0%; SCOFE 82;
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Pred. No.
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Pred. No.
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Pred. No.
            Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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            7.0%;
26.3%;
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26.3%;
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26.3%;
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Best Local Similarity 26.3%;
RESULT 1201
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US2002123463-A1.
05-SEP-2002.
(GETH ) GENENTECH INC.
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US2003032115-A1.
13-FEB-2003.
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(GETH ) GENENTECH INC.
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Best Local Similarity
                     Best Local Similarity RESULT 1197
                                                ABO09902 standard;
                                                                                                             Best Local Similarity RESULT 1198
                                                                                                                                        ABO08987 standard;
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Best Local Similarity
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23-JAN-2003.
                                                                                        23-JAN-2003.
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DB 6; Length 610;

44;

Query Match

Query Match

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Query Match

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Length 610;

DB 6;

44;

Length 610;

DB 6; 44;

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Length 610;

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DB 44;

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                      Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068752-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104539-A1.
                                                                                                                                                                                                    Human secreted/transmembrane protein (PRO) #46. US2003068685-A1.
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US2003068682-A1.
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Human secreted/transmembrane.protein (PRO) #46.
US2003073180-A1.
                                                                                                                                                                                                                                                                                                                                                                  Human secreted/transmembrane protein (PRO) #46 US2003068725-A1.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Local Similarity 26.3%;
                                   Query Match 7.0%;
Best Local Similarity 26.3%;
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26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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US2003068701-A1.
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US2003068755-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
     10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                    Query Match
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                                                                        RESULT 1216
                                                                                                                                                                                                    RESULT 1217
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                                                                                                   DB 6; Length 610; 44;
                                                                                                                                                                                                                                  Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 610;
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44;
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44;
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                                                                                                                                              ABU88528 standard; protein; 610 AA.
Human secreted and transmembrane polypeptide PRO872
US2002197615-A1.
26-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR95565 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054455-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR73973 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036135-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR80862 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049741-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABRB1167 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049743-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR88465 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068743-A1.
                                                                                                                                                                                                                                                                                                                                                                                              ABO05584 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032118-A1.
13-FBB-2003.
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44;
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44;
                               ABO08377 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO)
US2003044922-A1.
   26.3%; Pred. No. 44;
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Pred. No. 44;
                                                                                                   Score 82;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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26.3%; Pred. No.
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Human PR0872 polypeptide.
23-JAN-2003.
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26.3%;
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26.3%;
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RESULT 1212
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RESULT 1214
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1215
ID ABR8465 str
DE Humar
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                                                                                                                Best Local Similarity RESULT 1207
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RESULT 1210
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Best Local Similarity
RESULT 1211
Best Local Similarity
RESULT 1206
                                                                                  06-MAR-2003.
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Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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44;
                                                                                                         DB 6; Length 610;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064452-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR87550 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068705-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM77591 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054473-A1.
                                                                                                                                                                                                                                               Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                                                                                                              ABR94955 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003044930-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR95260 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003040075-A1.
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                                                              Human secreted/transmembrane protein (PRO) #46
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44;
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44;
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Human secreted/transmembrane protein (PRO)
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Pred. No. 44;
                                                                                                         Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                7.0%; Score 82; 26.3%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                      Score 82;
Pred. No.
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26.3%; Pred. No.
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                                                protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.0%;
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    7.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 1234
ID ABM27821 standard; J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1232
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               Best Local Similarity
RESULT 1225
                                                                                                                                                                                                                              Best Local Similarity
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                                                                                                                      Best Local Similarity RESULT 1226
                                                ABO03039 standard;
                                                                             US2003036131-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2003040071-A1.
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DB 6; Length 610; 44;
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44;
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                                          Length 610;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064447-A1.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064440-A1.
03-APR-2003.
                                                                        Human secreted polypeptide PRO872, SEQ ID NO:92. US2003068794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                         ABM26296 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104549-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABRÉ22820 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064462-A1.
                                                                                                                                                                                                                                                                                                         Human secreted polypeptide PRO872, SEQ ID NO:92.0S2003073183-A1.
                                                                                                                                                                                                  Human secreted polypeptide PRO872, SEQ ID NO:92 US2003068722-A1.
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01S2003065159-A1.
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44;
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Human secreted/transmembrane protein PRO872.
US2003008297-A1.
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                   protein; 610 AA
                                                                                                                                                                                         ABM03608 standard; protein; 610 AA
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17-APR-2003.
(GETH ) GENENTECH INC.
7.0%;
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Local Similarity .26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                           Best Local Similarity
RESULT 1236
                                                       Best Local Similarity
RESULT 1235
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Best Local Similarity
RESULT 1238
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                                                                                                                                                     Query Match
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Length 610;

DB 6; 44;

Score 82; Pred. No.

Length 610;

DB 6; 44;

Score 82; Pred. No.

6; Length 610;

DB 44;

Score 82; Pred. No.

Length 610;

9

DB 44;

Score 82; Pred. No.

Length 610;

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DB 44;

Score 82; Pred. No.

6; Length 610;

DB 44;

Score 82; Pred. No.

DB 6; Length 610; 44;

Score 82; Pred. No.

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DB 6; Length 610; 44;

Score 82; Pred. No.

6; Length 610;

DB 44:

Score 82; Pred. No.

Length 610

9

DB 44;

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ABM76371 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003082717-A1.
                                                                                                                                                                                                                                                                                                                             ABM76067 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104548-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM25991 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104543-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA21110 standard; protein; 610 AA.
Human secreted/transmembrane polypeptide PRO872.
US2003054404-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted/transmembrane protein (PRO) #46.
US2003036127-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO02429 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003040061-A1.
27-FEB-2003.
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                                                                         protein; 610 AA
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26.3%;
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26.3%;
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26.3%;
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Best Local Similarity 26.3%;
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26.3%;
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                                      26.3%;
                                                                                       Human PRO polypeptide #46.
US2003068732-A1.
       GENENTECH INC.
                                                                                                                          10-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                      Best Local Similarity RESULT 1254
                                                                                                                                                                                                                                                                                             Local Similarity
                     Query Match
Best Local Similarity
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RESULT 1261
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                                                                       ABO43671 standard;
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Best Local 8
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   (GETH )
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                                                        RESULT 1253
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                                      DB 6; Length 610;
                                                                                                                                                                        Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                               Length 610;
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44;
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                                                                                                                                                                                                                    ABMI5989 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064463-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM29041 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM07017 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068699-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM21111 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068707-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073175-A1.
                                                                                                   Human secreted polypeptide PRO872, SEQ ID NO:92 US2003073184-A1.
                                                                                                                                                                                                                                                                                                           DB 6;
44;
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US2003064451-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003068695-A1.
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44;
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44;
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44;
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Pred. No. 4
                                 Query Match 7.0%; Score 82;
Best Local Similarity 26.3%; Pred. No. RESULT 1244
                                                                                                                                                                                                                                                                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                  ABM02693 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                             ABO27550 standard; protein; 610 AA
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26.3%;
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RESULT 1250
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26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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US2003068703-A1.
                                                                                                                                   17-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1251
                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 1246
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RESULT 1252
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RESULT 1247

A B B B B

BEREE

RESULT

A D S E D

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6; Length 610;
                                                                                                                                                                                                                                                                                       7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
              Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Length 610; 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jocal Similarity 26.3%; Pred. No. 44;
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Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT.1268
ID ABR93430 standard; protein; 610 AA.
DE Human secreted polypeptide PRO872, SEQ ID NO:92.
PN US2003054478-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR71228 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003059880-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABRR 7855 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068718-A1.
                                                                                                                                                                                                            ABR04345 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003044917-A1.
                                                                                                                                                                                                                                                                                                                                         ABR75852 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003044929-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR93125 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003064465-A1.
                                                                  ABO16920 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003054470-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO27855 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003064454-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Apai0097 standard; protein; 610 AA.
Human secreted/transmembrane protein, PRO872.
US2003059831-A1.
                                                                                                                                                          Score 82; DB
Pred. No. 44;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                            7.0%; Score 82;
26.3%; Pred. No.
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26.3%;
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RESULT 1272
                7.0%;
                                                                                                                                                        Query Match 7.0%;
Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
                                                                                                                                           (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1266
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                          Best Local Similarity
RESULT 1263
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              Query Match
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DB 6; Length 610; 44;
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44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 610;
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44;
                                                                             Length 610;
                                                                                                                                                                                                      6; Length 610;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 610;
                                                                                                                                                                                                                                              ABM04887 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068727-A1.
                                                                                                                                                                                                                                                                                                                                                               ABMUNEW T Standard; protein; 610 AA. Human secreted polypeptide PROB72, SEQ ID NO:92. US2003068772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABMI0372 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003069407-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM11897 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104555-A1.
ABO29990 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003064461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein (PRO) #46.
US2003068714-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABO39497 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068776-A1.
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44;
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44;
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44;
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                             Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                7.0%; Score 82;
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Pred. No.
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Pred. No.
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                                                                                                                          protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO52043 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 7.0%;
Best Local Similarity 26.3%;
                                             PD 03-APR-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 26.3%;

RESULT 1273
                                                                                                                                                                       PD 10-APR-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCh

BEST Local Similarity 26.3%;

RESULT 1274
                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 26.3%;
RESULT 1275
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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RESULT 1277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polypeptide #46.
US2003068758-A1.
                                                                                                                       ABO33199 standard; protein
Human PRO polypeptide #46.
US2003068724-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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DB 6; Length 610; 44;

Length 610;

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DB (

Length 610;

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DB /

Length 610;

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Length 610;

DB 6; 44;

Length 610;

DB 6;

Length 610;

DB 6; 44;

Length 610;

DB 6;

Length 610;

DB 6;

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                            ABM06407 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068709-A1.
                                                                                                                                                                                     ABM04218 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068716-A1.
                                                                                                                                                                                                                                                                                                               ABM22331 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068740-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM07627 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003068751-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PRO872, SEQ ID NO:92 US2003073179-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM33127 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003087374-A1.
08-MAX-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein (PRO) #46. US2002068684-Al.
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                 7.0%; Score 82;
6.3%; Pred. No.
                                                                                                                                                                                                                                                                   Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                         7.0%;
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                                   26.3%;
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Human PRO polypeptide #46.
US2003049773-A1.
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ABM35364 standard, protein;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
    (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                          Query Match
Best Local Similarity
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                              Best Local Similarity RESULT 1291
                                                                                                              10-APR-2003
                                                                                                                                                                                                                                       10-APR-2003
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                 Query Match
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RESULT 1292
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                                                                                                                                                                                                                                                                                             Score 82; DB 6; Length 610;
Pred. No. 44;
                                                               Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Length 610;
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                                                                                                                                                                                       6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                         DB 6; Length 610;
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Pred. No. 44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054481-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR86940 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049778-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM28126 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054476-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM15252 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068692-A1.
                                                              9
                                                                                                                                                                                                                                ABO23666 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032134-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003068733-A1.
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44;
                                                              Score 82; DB
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 4
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                             ADA17641 standard; protein; 610 AA.
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RESULT 1290
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RESULT 1288
Human PRO polypeptide #46.
US2003049768-A1.
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US2003054987-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                         Best Local Similarity
RESULT 1282
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Best Local Similarity
RESULT 1283
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Best Local Similarity
RESULT 1286
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Best Local Similarity
RESULT 1284
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Best Local Similarity
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RESULT 1287

RESULT 1285

DB 6; Length 610; 44;

Length 610;

DB 6; 44;

Length 610;

DB 6; 44;

Length 610;

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Query Match

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Score 82; DB 6; Length 610;
Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003087373-A1.
                                                      ABM20806 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted polypeptide PRO872, SEQ ID NO:92 US2003073185-Al.
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US2003068773-A1.
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US2003951-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003049745-A1.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Local Similarity 26.3%; Pred. No.
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            Best Local Similarity 26.3%;
RESULT 1310
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Best Local Similarity 26.3%;
RESULT 1311
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO38582 standard;
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Pred. No. 44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM13422 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064457-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR80557 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049740-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049770-A1.
                                                                                                                                                                                                                                                                                                                                                           ABM18429 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054480-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003059885-A1.
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44;
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3 Human secreted/transmembrane protein (PRO) #46.

4 US2003036164-A1.

5 20-FBB-2003.

7.0%; Score 82; DB 6;
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RESULT 1302
ID ABOOSBB9 standard; protein; 610 AA.
DE Human secreted/transmembrane protein (PRO) #46.
PN US2003040074-A1.
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Human secreted/transmembrane protein PRO872
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44;
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44;
                                ABU99207 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO)
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44;
   26.3%; Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1303
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RESULT 1305
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RESULT 1309
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Best Local Similarity RESULT 1300
                                                                US2003040055-A1.
27-FEB-2003.
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Query Match

RESULT 1304

Query Match

RESULT 1306

Query Match

DB 6; Length 610;

Length 610;

9

DB 44;

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                    ABM28431 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003082715-A1.
                                                                                                                                                                                                                                                                                                                                       ABM28736 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003082716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM66180 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068737-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003096359-A1.
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                                                                                   Human secreted/transmembrane protein (PRO) #46. US2003068756-Al.
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Human secreted/transmembrane protein (PRO) #46.
US2003054454-A1.
20-MAR-2003.
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                                                 Score 82; DB
Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 1331
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Best Local Similarity
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Best Local Similarity
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    US2003068742-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1337
                                                                                                                                                                                            Length 610;
                                                                                                                                                                                                                                                                                                                          Length 610;
                                                                 Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 610; 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.0%; Score 82; DB 6; Length 610;
26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6; Length 610; 44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Length 610;
ABR96237 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003054458-A1.
                                                                                                          ABM02388 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003059886-A1.
                                                                                                                                                                                                                                      ABR86330 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003049758-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                                                                                                   ABR86635 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003049772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM29651 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM23856 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068735-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM23246 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068753-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM22026 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
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44;
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9
                                                            DB
44;
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44;
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Human secreted/transmembrane protein (PRO)
US2003068693-A1.
                                                                                                                                                                                            Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                       Score 82;
Pred. No.
                                                                 Score 82;
Pred. No.
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26.3%; Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%;
                                                               7.0%;
26.3%;
                                                                                                                                                     PD 27-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 26.3%;
RESULT 1321
                                                                                                                                                                                                                                                                                                                  Best Local Similarity 26.3%;
RESULT 1322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 26.3%; RESULT 1323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26.3%;
                                                                                                                                                                                                                                                                                        13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                    13-MAR-2003.
(GETH ) GENENTECH INC.
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                                                               Query Match
Best Local Similarity
RESULT 1320
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RESULT 1324
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RESULT 1325
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Length 610;

DB 6; 44;

Length 610;

DB 6; 44;

Length 610;

9

DB 44;

RESULT 1326

BBBBB

RESULT 1328

Length 610;

9 DB (

Length 610;

DB 6; 44;

A D M E I

Length 610;

DB 6; 44;

Length 610;

DB 6; 44;

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Query Match GENENTECH INC.
Query Match Best Local Similarity 26.3%;
RESULT 1348
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6; Length 610;
44;
                                                                                                                                                                                                                                Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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44;
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44;
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                                                                                                 6; Length 610;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068736-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM29346 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068679-A1.
                                                                                                                                                                                                                                                                                                                                                                                             ABR85720 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049753-A1.
                                                                                                                                             ABR96542 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054460-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM00558 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003073172-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR99702 standard, protein, 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM00253 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
                  ABO22108 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003054477-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                אפרביסיי (PRO) #46. US2003068700-Al. (PRO) #46. US2003068700-Al.
                                                                                                                                                                                                                                                                              ADA94129 standard; protein; 610 AA.
Human secreted/transmembrane protein PRO872
US2003059812-A1.
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44;
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44;
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Pred. No.
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                                                                                                 Score 82;
Pred. No.
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26.3%;
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                                                                20-MAR-2003.
(GETH ) GENENTECH INC.
Query Match
Pest Local Similarity 26.3%;
                                                                                                                                                                                                20-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1347
ID ABM29346 STD
PN Human
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RESULT 1344
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                                                                                                                                                                                                                                       Best Local Similarity
RESULT 1340
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RESULT 1338

ID ABO22100

DE Human se

PN US200300

PD 20-MAR-:

PA (GETH )
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DB 6; Length 610;
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44;
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44;
 Length 610;
                                                                                             Length 610;
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                                                                                                                                                                                                                           ABM20501 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104557-A1.
OS-UTW-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO16615 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003027276-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucroc41 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044920-A1.
 DB 6;
44;
                                                                                                                                                                                          DB 6;
44;
                                                                                                                                                                                                                                                                                                                          ADA81363 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003092121-A1.
                        Human secreted/transmembrane protein (PRO) #46. US2003068757-A1.
                                                                                             DB 6;
44;
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44;
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44;
Score 82;
Pred. No.
                                                                                             Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                          Match 7.0%; Score 82; Local Similarity 26.3%; Pred. No.
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Pred. No.
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US2003027265-Al.
OFER-2003.
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                                                                                                                        Human PRO polypeptide #46. US2003073182-A1.
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26.3%;
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RESULT 1355
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
                                                                      10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 1354
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US2003068719-A1.

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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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                                                                                        Length 610;
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26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 610;
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                                                                                                                                       ABM77896 standard, protein, 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
ABR81472 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                ABR89685 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073171-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted polypeptide PRO872, SEQ ID NO:92. US2003064458-Al.
                                                                                                                                                                                                                                                                                                                                                                                                        ABM26601 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032121-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM07322 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068702-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM01913 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068734-A1.
                                                                                    7.0%; Score 82; DB 6;
26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB030295 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003064464-A1.
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Human secreted/transmembrane protein (PRO) #46.
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26.3%; Pred. No. 44;
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Pred. No. 44;
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Pred. No.
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                            US2003049744-A1.
113-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                 Best Local Similarity
RESULT 1358
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RESULT 1359
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RESULT 1364
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Best Local Similarity
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Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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                                                                                                                                                                                                                   DB 6; Length 610;
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                                                                                                                                                                                                                                    Human secreted polypeptide PRO872, SEQ ID NO:92. 05-JUN-2003.
                                                                                                                                                                                                                                                                                                                             Human secreted/transmembrane protein (PRO) #46.
US200304942-A1.
                                                          ABO41632 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
                                                                                                                                                                                                                                                                                                                                                                                                                                               #46.
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Human secreted/transmembrane protein (PRO) #46.
US2003049750-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABOSOS18 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049779-A1.
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Human secreted/transmembrane protein (PRO)
US20030494747-A1.
13-MAR-2003.
(GETH ) GENENTECH INC.
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                      7.0%;
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26.3%;
                                                                                                               Best Local Similarity 26.3%;
RESULT 1368
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26.3%;
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Local Similarity 26.3%;
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RESULT 1375
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US2003049766-A1.
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US2003049767-A1.
10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1369
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Best Local Similarity
RESULT 1373
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Best Local Similarity
RESULT 1374
                                 Best Local Similarity
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Best Local Similarity
                                             T 1367
ABO41632 standard;
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                                                                                  US2003068729-A1.
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                       Query Match
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RESULT 1371
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US2003068696-A1.
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             Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                 DB 6; Length 610;
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Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                              7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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                                                                                                                                                  6; Length 610
                                                                                                                                                                                                  ABMI6904 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040078-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR96847 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054462-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064445-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM16294 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064449-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064441-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM14642 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                     ABR79642 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040059-A1.
                                                                                                                                                                                                                                                                                                                                                     ABO17936 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003044918-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA38554 standard; protein; 610 AA.
Human secreted/transmembrane protein PRO872.
US2003059780-A1.
                                                                                                                                              7.0%; Score 82; DB 26.3%; Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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               7.0%;
26.3%;
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RESULT 1381
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26.3%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 1382
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RESULT 1380
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                                Best Local Similarity
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                                                  RESULT 1376
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DB 6; Length 610;
                                                                                                                                                                                                                                                                                                                    Length 610;
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                                                Human secreted polypeptide PRO872, SEQ ID NO:92.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM75457 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104545-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US200310441-A1.
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US2003068775-A1.
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44;
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44;
                      Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 610 AA.
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                   Best Local Similarity 26.3%;
RESULT 1386
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RESULT 1387
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RESULT 1388
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Best Local Similarity 26.3%;
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Human PRO polypeptide #46.
US2003049765-Al.
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US2003049762-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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           GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                         ABO39192 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO46797 standard;
                                                                                                                                                                                                                                                                                                       17-APR-2003.
                                                                                                                                                                                                    10-APR-2003
10-APR-2003
(GETH ) GEN
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Query Match
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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                  DB 6; Length 610; 44;
                                                                                                                                                                                     DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                           DB 6; Length 610;
44;
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Pred. No. 44;
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44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6; Length 610
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                                                                                                                                     ABR71533 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032133-A1.
                                                                                                                                                                                                                ADM/2143 Standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003032136-A1.
                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PROB72, SEQ ID NO:92.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR73363 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054467-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054469-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR73058 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027270-A1.
06-FEB-2003.
                                              Human secreted/transmembrane protein (PRO) #46. US2003049752-A1.
                                                                                                                                                                                                                                                                                                                                                                                                 ABO06852 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 44;
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                   ADA83161 standard; protein; 610 AA.
                                                                                                 7.0%;
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26.3%;
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26.3%;
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26.3%;
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26.3%;
                                                                         13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
Best Local Similarity
RESULT 1395
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Best Local Similarity
RESULT 1404
                                                                                                 Query Match
Best Local Similarity
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                                      A D M E I
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Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
                                                                                                              DB 6; Length 610;
44;
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44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                ABR94040 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003059879-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR79947 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049738-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM11287 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064469-A1.
                                                ABO20583 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003032126-A1.
13-FEB-2003.
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Human secreted/transmembrane protein (PRO) #46.
US2003064466-A1.
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Human secreted/transmembrane protein PRO872.
US2003060407-A1.
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Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Human PRO polypeptide #46.
US2003054466-A1.
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US2003064453-A1.
                                                                                                                                                             protein; 610 AA
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    7.0%;
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Human PRO polypeptide #46.
US2003054463-A1.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                       Local Similarity
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Best Local Similarity
                                                                                                                           Best Local Similarity RESULT 1406
                   Local Similarity
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                     Best Loca
RESULT 1405
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Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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44;
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26.3%; Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068698-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068759-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM29956 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABMO5492 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003045700-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted/transmembrane protein (PRO) #46.
US2003068765-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003049748-A1.
Human secreted/transmembrane protein (PRO) #46.
US2003064468-A1.
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44;
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Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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13-Mnn - 13-
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10-APR-2003.
(GETH ) GENENTECH INC.
7.0%; S'
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Best Local Similarity 26.3%;
RESULT 1417
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 1420
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RESULT 1419
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RESULT 1421
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RESULT 1423
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                                                              03-APR-2003
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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                         Length 610;
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                                                                                                                                                                 Human secreted polypeptide PRO872, SEQ ID NO:92.05-JUN:2003.41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM26906 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068739-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABREBIGO standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068720-A1.
                                                                 Human secreted/transmembrane protein (PRO) #46.
US2003082767-Al.
                                                                                                                                                                                                                                                                                                           Human secreted/transmembrane protein (PRO) #46. US2003049774-Al. 13-MAR-2003. (SEM H) GENENTECH INC. 7.0%; Score 82; DB 6; Pt Local Similarity 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                 ABO49603 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049775-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA78456 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003073181-A1.
                                                                                                                                         DB 6;
44;
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Human secreted/transmembrane protein (PRO) #46
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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(GETH ) GENENTECH INC.
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                                                Best Local Similarity 26.3%;
RESULT 1424
                                       7.08;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1428
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
             10-APR-2003.
(GETH ) GENENTECH INC.
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US2003068688-A1.
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Length 610; Score 82; DB 6; Length 610; Pred. No. 44; Score 82; DB 7; Length 610; Pred. No. 44; Score 82; DB 7; Length 610; Pred. No. 44; DB 7; Length 610; 44; DB 7; Length 610; 44; 7; Length 610 7; Length 610; DB 7; Length 610; 44; 7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44; Human secreted polypeptide PROB72, SEQ ID NO:92. ABM17819 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040072-A1. ABR95870 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040073-A1. ABR77062 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92 US2003044927-A1. ABO49908 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003049776-A1. Human secreted/transmembrane protein (PRO) #46. US2003049780-A1. Human secreted/transmembrane protein (PRO) #46. US2003036126-Al. ABO21803 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003054475-A1. ABO24276 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003064467-A1. 7 ; ABO19973 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003032124-A1. DB 44; DB 44; DB 44; Score 82; Pred. No. Score 82; Pred. No. Score 82; Pred. No. Score 82; Pred. No. 7.0%; Score 82; 26.3%; Pred. No. 7.0%; Score 82; 26.3%; Pred. No. 7.0%; 7.0%; 7.0%; 7.0%; 26.3%; Query Match 7.0%; Best Local Similarity 26.3%; 7.0%; 7.0%; 26.3%; 13-MAR-2003. (GETH) GENENTECH INC. (GETH) GENENTECH INC. 13-MAR-2003. (GETH) GENENTECH INC. (GETH) GENENTECH INC. Query Match Best Local Similarity RESULT 1433 Best Local Similarity Query Match Best Local Similarity Best Local Similarity Best Local Similarity RESULT 1439 Best Local Similarity RESULT 1440 Best Local Similarity RESULT 1441 T 1436 ABR74583 standard; Best Local Similarity Best Local Similarity 06-MAR-2003. 06-MAR-2003. 20-MAR-2003 Query Match RESULT 1434 RESULT 8 2 2 2 2 2 E C D BE ID 2225

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Length 610;
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44;
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                                                                 ABR86025 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003049759-A1.
                                                                                                                                                                             ABM10677 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064455-A1.
                                                                                                                                                                                                                                                                                Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104550-A1.
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44;
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Pred. No. 4
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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US2003068728-A1.
03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Length 610;

DB 7; 44;

DB 7; Length 610; 44;

Query Match

Query Match

RESULT 1455

Query Match

RESULT 1453

Length 610;

7; DB 44; Length 610;

DB 44;

Length 610;

7;

DB 44;

DB 7; Length 610; 44;

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Query Match

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Human secreted polypeptide PRO872, SEQ ID NO:92.072803059894-Al.
                                                                                                                                                                                                                            мыкы7245 standard; protein; 610 AA.
Human secreted polypeptide PROB72, SEQ ID NO:92.
US2003068687-A1.
                                                                                                                                                                                                                                                                                                                                                                Human secreted polypeptide PRO872, SEQ ID NO:92. US2003303186-Al.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064443-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM24466 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064444-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068686-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM09762 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted/transmembrane protein (PRO) #46.
US2003068697-A1.
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Human secreted/transmembrane protein (PRO) #46.
US2003068710-A1.
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Pred. No.
                                                 Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%; Pred. No.
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Pred. No.
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Pred. No.
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Best Local Similarity 26.3%;
                                            Best Local Similarity 26.3%;
RESULT 1462
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RESULT 1464
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1 (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 26.3%;
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RESULT 1467
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Local Similarity 26.3%;
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PA (GETH ) GENENTECH INC.

Query Match 7.0%;

Best Local Similarity 26.3%;

RESULT 1465
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                27-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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US2003059882-A1.
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7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                 Score 82; DB 7; Length 610; Pred. No. 44;
                                                                                                                                                                                                                                                                            7; Length 610;
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44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036161-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR89075 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036119-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR72448 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036120-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR80252 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM01473 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                ABM19281 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104551-A1.
05-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR69036 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003027273-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO18546 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044921-A1.
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                                                                                                                                                                                                                                                                            DB 44;
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44;
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Human secreted/transmembrane protein (PRO)
US2003049757-A1.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Human PRO polypeptide #46.
US2003049761-Al.
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26.3%;
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            Best Local Similarity
RESULT 1452
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RESULT 1454
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RESULT 1459
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Best Local Similarity
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RESULT 1458
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RESULT 1460
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RESULT 1461
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Query Match

RESULT 1457

RESULT 1456

Query Match

DB 7; Length 610; 44;

DB 7; Length 610; 44;

Query Match

Query Match

Query Match

A B B B B B

7; Length 610;

DB 44;

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20-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
   US2003054457-A1.
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                                                 Query Match
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                               Length 610;
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Pred. No. 44;
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                                                                                                                                                      Score 82; DB 7; Length 610;
Pred. No. 44;
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44;
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44;
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44;
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                                                                                                                                                                              Human secreted polypeptide PRO872, SEQ ID NO:92. US2003104538-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR77667 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040067-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR78877 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054456-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
                    Query Match 7.0%; Score 82; DB 7;
Best Local Similarity 26.3%; Pred. No. 44;
RESULT 1471
                                                                          ABO38887 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003068774-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #46. US2003049781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein (PRO) #46
US2003036158-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 82; DB 7;
Pred. No. 44;
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US2002044806-Al.
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Human secreted/transmembrane protein (PRO)
US2003054482-A1.
                                                                                                                                                                                                                                                                                                                                                                                     Score 82;
Pred. No.
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Pred. No.
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US200330451-Al.
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26.3%;
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26.3%;
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17-APR-2003. (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1472
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Best Local Similarity
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Best Local Similarity
RESULT 1480
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2225

BEE

2882

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Length 610;
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   Length 610;
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                                                                                                                                                        Human secreted polypeptide PRO872, SEQ ID NO:92. US2003049764-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABMIJI17 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064450-A1.
                                                            Human secreted polypeptide PRO872, SEQ ID NO:92 US2003059883-A1.
                                                                                                                                                                                                                                                                                       ABR89990 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073177-A1.
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US2003064442-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068683-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM08237 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068754-A1.
DB 7;
44;
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44;
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Human secreted/transmembrane protein (PRO) #46.
US2003068731-A1.
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44;
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Pred. No. 44;
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                                                                                                                   DB 44;
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM27516 standard; protein; 610 AA
7.0%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1488
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               Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
                                              protein;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                              ABM01778 standard;
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7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
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Pred. No. 44;
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7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040065-A1.
                                                                                                                                                                                                            ABM74542 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003096351-A1.
                                                                                                                                                                                                                                                                                                                                                                                 ABM33737 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003096358-A1.
22-MAY-2003.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104556-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036122-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO46688 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049756-A1.
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Human secreted/transmembrane protein (PRO) #46.
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Human secreted/transmembrane protein (PRO) #46.
US2003036121-A1.
                                                           ABO40107 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003066681-A1.
10-APR-2003.
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Human secreted/transmembrane protein PRO872.
US2003017982-A1.
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RESULT 1492
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Best Local Similarity
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Best Local Similarity
RESULT 1496
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RESULT 1491
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RESULT 1494
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RESULT 1499
ID ABO15090 standard,
DE Human secreted/tran
               Best Local Similarity
RESULT 1490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-2003.
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Score 82; DB 7; Length 610; Pred. No. 44;
                                                                                                                                                                                             Score 82; DB 7; Length 610; Pred. No. 44;
PN US2003044919-A1.

PD 06-MAR-2003.

Query Match
Best Local Similarity 26.3%; Pred. No. 44;

RESULT 1500

ID AB017225 standard; protein; 610 AA.

DE Human secreted/transmembrane protein (PRO) #46.

PN US2003040077-A1.

PD 27-FRB-2003.

Query Match

7.0%; Score 82; DB 7; Les
                                                                                                                                                                                               7.0%;
                                                                                                                                                                                                                   Local Similarity
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Length 229;
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(EGET ) GENENTECH INC.
PLY MALCh TG.0%; Score 174; DB 3; Length 229;
Ft Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB44310 standard; protein; 229 AA.
Human PRO733 (UNQ411) protein sequence SEQ ID NO:447.
WO200053756-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB34734 standard; protein; 229 AA.
Human secreted protein encoded by DNA clone vg20 1.
WO200055375-A1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US4001-1-0.
29-JUL-2004.
29-JUL-2004.
100.04; Score 229; DB 8; I
ery Match 100.04; Score 229; DB 8; I
                                                                                                                                                                                         PD 23-OCT-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 229; DB 8; I Best Local Similarity 100.0%; Pred. No. 1.5e-211; RESULT 8
                                                                                               100.0%; Score 229; DB 7; 100.0%; Pred. No. 1.5e-211;
                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 229; DB 8; 100.0%; Pred. No. 1.5e-211;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human p24 vesicle trafficking protein Hp24-2.
WO9836068-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.0%; Score 174;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                       ADL91839 standard; protein; 229 AA.
Human PRO83480 protein SEQ ID NO:60
WO2004024076-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 229 AA.
                                                                                                                                                        protein; 229 AA.
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Human protein clone HP02515.
WO200005367-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW49100 standard; protein; 229 AA
                                                                                                                                                                            Human T1R-like ligand II protein. US2003198618-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human T1R-like ligand II protein.
US2004146501-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY41754 standard, protein; 22
Human PRO733 protein sequence.
WO9946281-A2.
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(SAGA ) SAGAMI CHEM RES CENT.
(PROT-) PROTEGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
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(ALPH-) ALPHAGENE INC.
76.0%;
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(INCY-) INCYTE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                               25-MAR-2004.
(GETH ) GENENTECH INC.
                   29-MAY-2003.
(NIJJ/) NI J.
(GENT/) GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Best Local Similarity
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     US2003100048-A1.
                                                                                                 Query Match
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                                       (without alignments)
541.340 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES
                                                                                                               1 MGDKIWLPFPVLLLAALPPV.....SAIQVYMLKSLFEDKRKSRT 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description
OM protein - protein search, using sw model
Run on: March 19, 2007, 23:13:05 ; Search time 207 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; SCI INC.
100.0%; Score 229; DB 2; Length 229;
100.0%; Pred. No. 1.5e-211;
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(HUMA-) HUMAN GENOME SCI INC.
ery Match
ery Match
ery match 100.0%; Score 229; DB 4; Length 229;
ery match 100.0%; Pred. No. 1.5e-211;
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(EOSB-) EOS BIOTECHNOLOGY INC.
ery Match
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incomplexity 100.0%; Pred. No. 1.5e-211;
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ry Match 100.0%; Score 229; DB 7; Length 229;
t Local Similarity 100.0%; Pred. No. 1.5e-211;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB70497 standard; protein; 229 AA.
Angiogenesis protein AAA4 protein sequence (Fig 4)
WO200111086-A2.
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US6586210-Bl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 229; DB 6; 100.0%; Pred. No. 1.5e-211;
                                                                                                                                              Gapop 60.0 , Gapext 60.0 2782304 seqs, 489333398 residues Total number of hits satisfying chosen parameters: Minimum DB seq length: 2000000000
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Human T1-receptor-(T1R-)like ligand II.
US2002142461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW48335 standard; protein; 229 AA.
Human T1 receptor-like ligand II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAE02806 standard; protein; 229 AA. Human TIR-like ligand II protein. WO200142506-A1.
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Human T1 receptor-like ligand II.
                                                                                                                                                                                                                                                                                                                                                                                 geneseqp2003bs:*
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i: geneseqp1980s:*
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GENTZ R L

JEE/) RUBEN S M.

JET/ Match

Best Local Similarity

RESULT 5

ID ADB87500 stand

DE Human T1 r

PN US65867

PD 01-
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(HUMA-) HUMAN GENOME
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Best Local Similarity
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                                                                                             Perfect score:
                                                                                                                                 Scoring table:
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Human secreted/transmembrane protein (PRO) #205.
US2003036179-A1.
                                                                    20-FEB-2003
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 6; Length 229;
ery Match 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABU72262 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2002192706-A1.
                                                                                                                                                                                                                                ABO17820 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003032156-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO733.
US203050239-A1.
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Human secreted and transmembrane PRO polypeptide #18.
US2002177553-A1.
                                                                                                                                                                                                                                                                                                                U3-FEB-2003.
(GETH ) GENENTECH INC.
Fry Match 76.0%; Score 174; DB 6; Leery Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
76.0%; Score 174; DB 6; LE.
iry Match.
iry Match. 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 4; Lk
ery match 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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Novel secreted and transmembrane protein PRO733.
US2003017563-A1.
     AAU12376 standard; protein; 229 AA.
Human PRO733 polypeptide seguence.
WO200140466-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABU66774 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003036180-A1.
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Human PRO733 polypeptide.
US2002169284-A1.
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ID AB025045 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU81074 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S2003cc__
20-FBB-2003.
(GETH ) GENENTECH INC.
76.0%; Sc
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US2003004311-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-2002
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Length 229;
                                                                                                                                                                            76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
17 Match 76.0%; Score 174; DB 6; Length 229;
8t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                   ABUG7050 standard; protein; 229 AA.
Human secreted/transmembrane, PRO, protein SEQ ID 410.
US2003032155-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA45929 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003022328-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB19418 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003068796-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2005.
(GETH) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 6; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
FRY MATCh 76.0%; Score 174; DB 6; LK
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Pred. No. 1.2e-158;
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CGETH GENERIECH INC.

17 Match

76.0%; Score 174; DB 6; L

17 Match

17 Theal Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                                ABG72289 standard; protein; 229 AA.
Human p24 vesicle trafficking protein Hp24-2.
US2002132999-A1.
                                                                               ABU80409 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733.
US2003004102-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA19010 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003054517-Al.
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Human PRO polypeptide #205.
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17-APR-2003.
(GETH ) GENENTECH INC.
76.0%; Scr
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) GENENTECH INC.
76.0%; SCC
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19-SEP-2002.
(INCY-) INCYTE PHARM INC.
Orv Match -: -: 1avity 100.0%;
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(GETH ) GENENTECH INC. 76.0%;
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(GETH ) GENENTECH INC.
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US2003073212-A1.
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RESULT 26
                              Query Match
Best Local Similarity
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US2003049816-Al.
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Length 229;

Length 229;

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Score 174; DB 6; Length 229;
Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Ouery Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                    мымулыз5 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel human secreted and transmembrane protein PRO733.06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA24986 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003050241-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human secreted and transmembrane protein PRO733. US2003082691-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           мылывы standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003073211-A1.
               A (GETH) GENENTECH INC.
Query Match
76.0%; Score 174; DB 6; L.
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                           Score 174; DB 6; Lo
Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.

FOR MATCH 76.0%; Score 174; DB 6; LA

FOR MATCH 76.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                               ADB16743 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                  ADB14898 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087351-A1.
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1925003077722-A1.
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US200082710-A1.
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08-MAY-2003.
(GETH ) GENENTECH INC.
76.0%; SC
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Query Match
76.0%; 8
Best Local Similarity 100.0%;
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24-APR-2003
(GETH ) GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC. 76.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 51
   08-MAY-2003
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(GETH ) GENENTECH INC.

(GFTH ) GENENTECH INC.

(GFT) Match 76.0%; Score 174; DB 6; Length 229;

(FT) Timilarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(Match 76.0%; Score 174; DB 6; Length 229;
+ 1.0cal Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                OG-MAY-2003.
(GETH ) GENENTECH INC.
ery Match
76.0%; Score 174; DB 6; Length 229;
ery Match
76.0%; Pred. No. 1.2e-158;
                                                        DB 6; Length 229;
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                                                                                               ALMANASA Standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733
US203082111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA85886 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA87541 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087345-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Uszuszuszus
Un-APR-2003.
(GETH) GENENTECH INC.
FLY Match 76.0%; Score 174; DB 6; Le
ETY Match 100.0%; Pred. No. 1.2e-158;
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10-APR-2003.
(GETH) GENENTECH INC.
FLY Match 76.0%; Score 174; DB 6; Le
FLY Match 100.0%; Pred. No. 1.2e-158;
                                                      76.0%; Score 174; DB 6; Lot 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 6; Lk
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100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                     Lucas scandard; protein; 229 AA. Human PRO polypeptide #205.
US2003087350-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #205.
US200307315-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA67583 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003068795-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB30590 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003068794-A1.
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Human PRO polypeptide #205.
US2003082705-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA79402 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003082763-A1.
                 01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 40
                                                                                                                                                                                                                      Best Local Similarity RESULT 35
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Best Local Similarity
RESULT 39
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                                                                        Best Local Similarity
US2003082704-A1.
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                                                        Query Match
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Length 229;

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PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 6; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 65
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(GETH) GENENTECH INC.
ery Match 76.0%; Score 174; DB 6; Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

ery Match 76.0%; Score 174; DB 6; Length 229;

ery Match 76.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 6; Length 229;
                                                                                                                                                                   ID ADA75808 standard; protein; 229 AA.

BE Human PRO polypeptide #205.

PN US2003082703-A1.

PD 01-MAY-2003.

Query Match

Query Match

RESULT 63

RESULT 63
                                                                                                                       (GETH) GENENTECH INC.

ry Match 76.0%; Score 174; DB 6; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
76.0%; Score 174; DB 6; Length 229;
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 76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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24-TR-2003.
24-TR-2003.
GETH ) GENENTECH INC.
76.0%; Score 174; DB 6; LA
ery Match
76.0%; Score 174; DB 6; LA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide SEQ ID NO 410. 24.280.007715-A1.
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US200307774-A1.
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15-May - 1.05.003092147-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB31142 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003096386-A1.
                                                                                                                                                                                                                                                                                                                                       ADA47033 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003073210-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA93505 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003077721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA61070 standard; protein; 229 AA
                                                     protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                     USACOSO
17-APR-2003
(GETH ) GENENTECH INC.
76.0%;
                                                   ADA80566 standard; protein;
Human PRO polypeptide #205.
US2003082761-A1.
 Query Match
Best Local Similarity
RESULT 61
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Best Local Similarity
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Best Local Similarity
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US2003049817-A1.
                                                                                                                                             Query Match
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(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

76.0%; Score 174; DB 6; Length 229;

ery Match

76.0%; Pred. No. 1.2e-158;
                  76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                          76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                          Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003082708-A1.
                                                                    ABO19711 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003050240-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA85334 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003082695-A1.
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USACCOLORS.
(GETH ) GENENTECH INC.
16ETV Match 76.0%; Score 174; DB 6; LEV Match 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                PD 20-MAR-2003.
PA (EETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                              76.0%; Score 174; DB 6; L
100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                Human secreted/transmembrane polypeptide PRO733 US2003055216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB24769 standard; protein; 229 AA. Human PRO polypeptide SEQ ID NO 410 US20030777713-A1.
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01-Mav-002701-A1.
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US2003073214-A1.
                                                                                                                                                                                                                                                                                                                                                         ADA74536 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003068798-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA75256 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003073216-A1.
                                                                                                                                                                                                                 ADA12647 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 55
                                                                                                                                                                         Best Local Similarity RESULT 53
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         Query Match
Best Local Similarity
RESULT 52
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ADA88093 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082700-A1.
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US2003082686-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADA97650 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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                                                                                                                                                    Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 78
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO19602 standard; protein; 229 AA.
Novel human secreted and transmembrane polypeptide #70
US2003049633-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB21788 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082765-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003082709-A1.
                                                                                                                                                                                                                                                                                                             76.0%; Score 174; DB 6; Lo
100.0%; Pred. No. 1.2e-158;
        100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.0%; Score 174; DB 6; I 100.0%; Pred. No. 1.2e-158;
                                            ADA96546 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082690-A1.
                                                                                                                                                                                                           ADA81118 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082702-A1.
                                                                                                                                                                                                                                                                                                                                                                        ADA95994 standard, protein, 229 AA. Human PRO polypeptide #205.
US2003082759-Al.
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Human PRO polypeptide #205.
US2003082760-A1.
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0S200307710-A1.
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Human PRO polypeptide #205.
US2003068797-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                               Best Local Similarity RESULT 71
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Best Local Similarity RESULT 70
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RESULT 76

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20-MAR-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
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(GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
0 0-MAY-2003.

4 (GETH) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                            ADA88645 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003073213-A1.
17-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB22340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087344-A1.
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(GETH ) GENENTECH INC.
Sry Match 76.0%; Score 174; DB 7; Le
ترميماً Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Beet Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 86
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01-mav-non
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1052003059909-Al.
                                                                                                                                                                                                                         ADB28511 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082699-A1.
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Human PRO polypeptide #205.
US2003022239-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA67031 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003068793-A1.
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27-MAR-2003.
(GETH ) GENENTECH INC.
76.0%; SCC
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17-APR-2003.
(GETH ) GENENTECH INC.
76.0%; Scc
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(GETH ) GENENTECH INC.
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24-APR-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
                                                                              76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2003.
(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                     Novel human secreted and transmembrane protein PRO733. US2003082687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB77370 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082696-A1.
                                                                                                                                    ADB39535 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082764-A1.
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(GETH) GENENTECH INC.

(GETH) MAtch

(ery Match

(amilarity 100.0%; Pred. No. 1.2e-158;
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US2003077716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB34527 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077717-A1.
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                                                                                                                                                                                                                                                                         ADB73953 standard; protein; 229 AA. Human PRO polypeptide #70. US2003045462-A1. GG-MAR-2003. (GETH ) GENENTECH INC. PF.0%; Score ery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 229 AA
                                                 (GETH ) GENENTECH INC.
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US2003083248-Al.
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(GETH ) GENENTECH INC.
                                                                                              Best Local Similarity RESULT 98
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                       Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003082712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB38702 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003082766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB38150 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087347-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 we secreted and transmembrane protein PRC to 01-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158; ID ADB38150 standard; protein; 229 AA.

PN USZ003087347-A1.

PA (GETH ) GENENTECH INC.

76.0%; Score 174; DB 7; Leng RESULT 99 PN USZ003087347-A1.
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PA (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; DB 7; L

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                   Score 174; DB 7; L
Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
76.0%; Score 174; DB 7;
76.0%;
                                                                                                                                                                                                                                                                                  Human PRO polypeptide SEQ ID NO 410. 125.08307712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB15450 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB90434 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082762-A1.
01-MAY-2003.
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24-APR-2003,
(GETH ) GENENTECH INC,
Match 76.0%; Sco
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RESULT 95

RESULT 92

Length 229;

Length 229;

us-10-692-730-2.olig30_spdi.rag

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Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein Seg ID410.
US2003087365-A1.
   Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                              Best Local Similarity
RESULT 118
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                      RESULT 116
                                                                                                                                                                    RESULT 117
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                                           BEREE
76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                     AUM46578 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2002082692-A1.
                                                                                                42-APR-2003.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 7; Le
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24-APR-2003.

26-APR-2003.

76.0%; Score 174; DB 7; Le

16-158; Score 174; DB 7; Le
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Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC44095 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003054986-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC61855 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003049684-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC66919 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003060406-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC63819 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003054405-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC63103 standard, protein, 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003068648-A1.
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(GETH ) GENENTECH INC.
76.0%; SCOIE 174;
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                                                 ADB35079 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077718-A1.
                                                                                                                                                                                                   ADB36183 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077720-A1.
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01-MAY-2003.
(GETH ) GENENTECH INC. 76.0%; S
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
             Best Local Similarity RESULT 107
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Best Local Similarity
RESULT 109
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Best Local Similarity
RESULT 108
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Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003092106-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC71998 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092107-A1.
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Novel human secreted and transmembrane protein PRO733
US2003092105-A1.
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(GETH) GENENTECH INC.
(GETH) GENENTECH INC.
(ery Match
(ery Match) 76.0%; Score 174; DB 7; Le
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(GETH ) GENENTECH INC.
sry Match 76.0%; Score 174; DB 7; Le
''---1 similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; Lesty Match 76.0%; Pred. No. 1.2e-158;
                                                                      10-APR-2003.
A (GETH) GENENTECH INC.
OLORY MALCh
Pest Local Similarity 100.0%; Pred. No. 1.2e-158;
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(EGTH) GENENTECH INC.
176.0%; Score 174; DB 7; Le
st Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                   76.0%; Score 174; DB 7; I 100.0%; Pred. No. 1.2e-158;
ADC68168 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003069178-A1.
                                                                                                                                                                                                            Human secreted/transmembrane protein, PRO733 US2003072745-A1.
                                                                                                                                                                                                                                                                                                                                                                 ADC67543 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003073131-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC62479 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003073624-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC42112 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
                                                                                                                                                                                                                                                                                                                                                                                                                                (GETH) GENENTECH INC.
76.0%; Score 174;
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                                                                                                                                                                                                                                             17-APR-2003.
(GETH ) GENENTECH INC: 76.0%;
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) GENENTECH INC.
76.0%;
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protein; 229 AA

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Novel human secreted and transmembrane protein PRO733. US2003092104-A1.
  ADD03235 standard;
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                                                                                                                                                                                                                                                                             Length 229;
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    AUC5/338 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seq ID410.
US2003087366-A1.
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Novel human secreted and transmembrane protein Seg ID410.
US2003087364-A1.
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Novel human secreted and transmembrane protein Seg ID410.
US2003087359-A1.
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Novel human secreted and transmembrane protein Seg ID410.
US2003087360-A1.
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Novel human secreted and transmembrane protein Seg ID410.
US2003087346-A1.
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Novel human secreted and transmembrane protein Seg ID410.
US2003087363-A1.
                                                                                                                                                                           ADC60529 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                                                       ADC51004 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087361-A1.
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US-MAY-2003.

(GETH) GENENTECH INC.

(GETY MATCh 76.0%; Score 174; DB 7; Le

(ery Match 76.0%; Pred. No. 1.2e-158;
                                                                     OB-MAY-2003.
(GETH ) GENENTECH INC.
T6.0%; Score 174; DB 7; Le
lery Match 76.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                         76.0%; Score 174; DB 7; Lot 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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US2003087362-A1.
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100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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RESULT 125
1D ADC573
DE Novel
PN US2003
PD 08-MAY
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PD 15-MAY-2003.

PA (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; 'DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
st_iocal Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 136
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 229;
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Pred. No. 1.2e-158;
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Query Match 76.0%; Score 174; DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ry Match
Total Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                               ADC90227 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087348-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD04639 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC80595 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003092103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                  Score 174; DB 7; Li
Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                           Human PRO polypeptide #205.
1052003194770-A1.
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1052003194773-A1.
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Human PRO polypeptide #205.
US2003194776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADD11102 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003194774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC47983 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003194771-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC80043 standard; protein; 229 AA
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16-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; S
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(GETH ) GENENTECH INC.
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GENENTECH INC.
76.0%;
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Length 229;

Length 229;

Length 229;

Length 229;

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76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE32351 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-OCT-2003.
(GETH ) GENENTECH INC.
rry Match 76.0%; Score 174; DB 7; Lerry Match 76.0%; Pred. No. 1.2e-158;
                                                                                                                                                                 ) 22-MAY-2003.

(GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 7; L

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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( GETH ) GENENTECH INC.

QUETY MAICh 76.0%; Score 174; DB 7; L.

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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A (GETH) GENENTECH INC.
OLORY MATCH
Dest Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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( (GETH) GENENTECH INC.

76.0%; Score 174; DB 7; L

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003203434-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB16649 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203435-A1.
                                                                                                                                   Human secreted/transmembrane protein, PRO733 US2003096744-A1.
                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #205.
US2003199055-A1.
                                                                                                                                                                                                                                                             ADD92544 standard; protein; 229 AA Human PRO polypeptide #205.
US2003199030-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE22283 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003199056-A1.
                                                                                                                ADE49481 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 229 AA.
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(GETH ) GENENTECH INC.
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Human PRO polypeptide #205.
US2003199057-A1.
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US2003203428-A1.
                      30-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD79507 standard;
       US2003203432-A1.
                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                           RESULT 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 156
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                               GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

ery Match

76.0%; Score 174; DB 7; Length 229;
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1.2e-158;
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                 ADD1225 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733 US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD53656 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003203437-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD54227 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                                                    30-OCT-ZUUS.
(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
iry Match 76.0%; Score 174; DB 7; Le
iry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
76.0%; SCOIE 174;
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Pred. No.
                                                                                                                         ADD09512 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003194775-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                         ADD52364 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194769-AI.
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Human PRO polypeptide #205.
US2003194779-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD02611 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003203431-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD02045 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003203430-Al.
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Human PRO polypeptide #205.
US2003194792-A1.
16-OCT-2003.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
RESULT 147
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                                                                                    Best Local Similarity RESULT 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 149
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RESULT 148
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Best Local Similarity
RESULT 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                    US2003087358-A1.
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                                                                           Query Match
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ID ADDS42
DE Novel
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PD 23-OCT-2003.

PD A (GETH ) GENENTECH INC.

Query Match

Auch

Rest Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 172

ID ADE18964 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OSCT_2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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AA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
                                Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE32903 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194766-A1.
                                                                                                                                                                                                                                                                                                                               Luciy match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD95949 standard; protein; 229 AA.

Human PRO polypeptide #205.
US2001199059-A1.
US2001199059-A1.
(GETH) GENENTECH INC.
T6.0%; Score 174; DB 7; L.

Stry Match
T6.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) 30-OCT-2003.

1 (GETH) GENENTECH INC.

76.0%; Score 174; DB 7; L

Dest Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane protein, PRO733.
US2002203433-Al.
                                                                                                                                                                                                                                                                                                                                                                               Human PRO polypeptide #205.
US2003199033-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #205.
US2003199064-A1.
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30-000-001
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US2003199026-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE42595 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199032-A1.
                                                                                         ADE19516 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199025-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                  23-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                            Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.0%; Score 174; DB 7; Length 229; RESULT 169
                                                    Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                         Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE33455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE34007 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194791-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) 23-OCT-2003.
A (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 7; L
1.2e-158;
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(GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD72622 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003194781-A1.
                                                                                                            ADD72264 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-NOV-2003.
(GETH ) GENENTECH INC.
76.0%; Score 174;
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Best Local Similarity 100.0%; Pred. No.
RESULT 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO polypeptide #205.
US2003199053-Al.
                                                                                                                                                                                                                                                   Luman PRO polypeptide #205.
US2003194772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                ADE17860 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD80059 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003207417-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD33096 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194768-A1.
                                                                                                                                                                                                                                                                                                                       16-OCT-2003. GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                 30-OCT-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 168
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Length 229;

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76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                            76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 186
                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 7; Length 229;
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(GETH ) GENENTECH INC.
76.0%; Score 174; DB 7; Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207355-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG23201 standard, protein, 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207384-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-CCT-2003.
(GETH) GENENTECH INC.
(ETY Match 76.0%; Score 174; DB 7; Leery Match 76.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 76.0%; Score 174; DB 7; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF47287 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195333-A1.
                                                      ADD80611 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207418-A1. GG-NOV-2003. GGFTH ) GENENTECH INC.
                                                                                                                                                                                             ADD89639 standard; protein; 229 AA. Human PRO polypeptide #205. US2003199028-AI. GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #205.
US2003199034-Al.
                                                                                                                                                                                                                                                                                                                       RESULT 182
ID ADE40923 standard; protein; 229 AA.
DE Human PRO polypeptide #205.
PN US2003199031-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE92851 standard; protein; 229 AA
Human PRO polypeptide #205.
US2003194777-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE95520 standard; protein; 229 AA. Human NOVX4c protein. WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(CURA-) CURAGEN CORP.
Query Match
Best Local Similarity
RESULT 180
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RESULT 181
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PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 190
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Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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16.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 196
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(GETH ) GENENTECH INC.
ry Match
    76.0%; Score 174; DB 7; Length 229;
    7-ran Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733
US2003207379-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH55340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733 US2003207381-A1.
                                                                                                                                                                                                                                                                                                                        06-NOV-zuvs.
(GETH ) GENENTECH INC.
rry Match 76.0%; Score 174; DB 7; Le
rrann Similarity 100.0%; Pred. No. 1.2e-158;
Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GETH) GENENTECH INC.

ry Match
76.0%; Score 174; DB 7; Lv
rocal Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Lust Local Similarity 100.0%; Pred. No. 1.2e-158;
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Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG51044 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003216561-A1.
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Human secreted/transmembrane protein, PRO733.
US2003206915-A1.
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US2003077700-Al.
                                                     ADF97536 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003207370-A1.
                                                                                                                                                                                                                                           ADG80600 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207373-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG80048 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207372-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 192
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ADC81147 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092115-A1.
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
ery Match 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 206
                                                                                                                                                                                                                                                                                               Length 229;
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                                                                                                                                                                     AU1031559 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US203207387-A1.
                 AD165060 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207386-A1.
O6-NOV-2003.
(GETH ) GENENTECH INC.
Fry Match
St. Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003087385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADNI5437 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087356-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADN14885 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087357-A1.
                                                                                                                                                                                                                                                                                                                                                       ADH81973 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADH81421 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207377-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM82590 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087355-A1.
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Novel human secreted and transmembrane protein PRO733
US2003087353-A1.
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(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
:ry Match 76.0%; Score 174; DB 7; Le
ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                         Best Local Similarity RESULT 199
                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 200
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RESULT 203
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                                                                                                                                                                                                                                                                                                                                                                                                  US2003207388-A1.
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                                                                                                                            Query Match
  RESULT 198
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15-MAY-2003.
(GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
ery Match
-- ' ' And Similarity 100.0%; Pred. No. 1.2e-158;
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ry Match
76.0%; Score 174; DB 8; Length 229;
t Local Similarity 100.0%; Pred. No. 1.2e-158;
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 208
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 76.0%; Score 174; DB 8; Length 229; Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
15-08; Score 174; DB 8; Length 229; Sty Match
St Local Similarity 100.08; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          / Match 76.0%; Score 174; DB 8; Length 229; Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 229;
                                                                                                                                                                                                                                                                                                                                                                        15-MAY-2003.
(GETH) GENENTECH INC.
176.0%; Score 174; DB 8; L.
St Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003104536-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #205.
102003203440-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE75811 standard; protein; 229 AA. Human PRO polypeptide #205. US2003211571-A1.
                                                                                                                      ADD76595 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003100087-A1.
                                                                                                                                                                                                                                                                                                  ADD87959 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003092113-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE23387 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003092108-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE23939 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003092110-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE24582 standard, protein, 229 AA. Human PRO polypeptide #205.
US2003092111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD87407 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003zcj.
30-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; SCO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-Zuus.
(GETH ) GENENTECH INC.
76.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 209
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(GETH ) GENENTECH INC.
ry Match 76.0%; Score 174; DB 8; Length 229;
t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                        PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 223
                                                                                                                                                                                                   16-OCT-2003.

(GETH) GENENTECH INC.

Query Match

16.0%; Score 174; DB 8; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PD 23-OCT-2003.

PA (GETH.) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 228
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                               Length 229;
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(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 8; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; LA

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 23-OCT-2003.

(GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; L.

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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A (GETH) GENENTECH INC.

76.0%; Score 174; DB 8; L.

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
   ADF40214 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003198994-A1.
                                                                                                                                            ADF46010 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195148-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, PRO733. US2003203402-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APF24406 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003204055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF40838 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199021-A1.
                                                                                                                                                                                                                                                                                     ADE94741 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199027-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                ADE91152 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE93403 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003199060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF33765 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE95293 standard; protein;
Human PRO polypeptide #205.
US2003199052-A1.
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                PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 8; Length 229;
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                                                                                                                                                           PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; LA Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 218
                                                                                                                                                                                                                                                                                                      PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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A (GETH ) GENENTECH INC.
OLORY MAICH
Dest Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003130181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF61522 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195345-A1.
                                                                                                                                                                                                                                                    Human PRO polypeptide #205.
1052003194794-Al.
                                                                                                                        ADE89273 standard; protein; 229 AA
Human PRO polypeptide #205.
US2003199062-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          ADE88721 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003199054-A1.
Human PRO polypeptide #205.
US2003203439-A1.
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BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PANJ) PAN J. (PANJ) PAN J. (ROYM) PANN N. (SHEL) SHELTON D L. (STEM) STEMART T A. (TUMA) TUMAS D. (WILL) WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GODOWSKI P J. GIRMALDI J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DESNOYERS L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILVAROFF E
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KLJAVIN I J
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Best Local Similarity
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NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GAO W.
GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODDARD A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GURNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FONG S.
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(BAKE/)
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Query Match
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              PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                              Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUSY2299 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003199051-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-ZUUS.
(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 8; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Le
Query Match 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                               US.C.T. 2003.
(GETH ) GENENTECH INC.
16 Y Match 76.0%; Score 174; DB 8; L. 16 Y Match 75.0%; Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76.0%; Score 174; DB 8; L Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             76.0%; Score 174; DB 8; L
100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003199435-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF26608 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
Human secreted/transmembrane protein, PRO733. US2003194780-A1.
                                                                                                                                                                                                                                                                                      ADF27232 standard; protein; 229 AA.:
Human secreted/transmembrane protein, PRO733.
US2003199436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF27868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199437-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF33141 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003211091-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF25507 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003211092-A1.
                                                                                                                                    ADF34984 standard; protein; 229 AA. Human PRO polypeptide #205. US2003199029-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE90600 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO polypeptide #205.
US2003199063-A1.
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2003.
214 ) GENENTECH IN.
217 Match
Best Local Similarity 1
RESULT 234
ID ADE27868 standa-
DE Human secre-
PN US20031
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENENTECH INC.
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(GETH) GENENTECH INC. 76.0%; Score 174; DB 8; Length 229; ry Match 100.0%; Pred. No. 1.2e-158;
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                                                                                Length 229;
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(GETH ) GENENTECH INC.
PLY MALCH
Ft Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207426-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE91747 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003199058-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG22112 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207360-A1.
US2003199674-A1.
23-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; Le
                                                                                                                                                                                        OCCT-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Luery Match 76.0%; Pred. No. 1.2e-158;
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Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
PY MATCH 76.0%; Score 174; DB 8; Lust Match 100:0%; Pred. No. 1.2e-158;
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A (GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; L.
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158
                                                                                                                       Human secreted/transmembrane protein, PRO733. US2003194410-A1.
                                                                                                                                                                                                                                                                                          ADF46634 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003195344-A1.
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US2003207422-A1.
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Human PRO polypeptide #205.
US2003207376-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG02326 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF98659 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003208055-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC.
76.0%; S
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                                                                                              Best Local Similarity
RESULT 241
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Best Local Similarity
RESULT 242
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DB 8; Length 229;

DB 8; Length 229;

Length 229;

Length 229;

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OG-NOV-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
                 (GETH ) GENENTECH INC.
179 Match
181 Ancal Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(GY Match 76.0%; Score 174; DB 8; Length 229;
(ery Match 76.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO733.
US2003207427-A1.
                                                                                                                                                                                                                                                 ADG23753 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207389-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human secreted and transmembrane protein PRO733 US2003207350-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207428-A1.
(GE-NOV-2007)
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG07792 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207356-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG55287 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003194778-A1.
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(GETH ) GENENTECH INC.
ery Match
16.0%; Score 174; DB 8; List Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 264
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 263
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16-OCT-2003.
(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 8; L.
                                                                                                 ADG06169 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207374-A1.
                                                                                                                                                                                                                                                                                                                                                                                                 ADG04042 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207423-A1.
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                                                                                                                                                                                                                                                                                                       06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                 Best Local Similarity
RESULT 259
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Best Local Similarity
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                                            Query Match
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PD 06-NOV-2003.
A (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 250
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(GETH ) GENENTECH INC.
1017 Match 76.0%; Score 174; DB 8; Length 229;
                                                                                                                                                                                                               76.0%; Score 174; DB 8; Length 229; 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 256 Similarity 100.0%; Pred. No. 1.2e-158;
TD ADG08416 standard; protein; 229 AA.
DE Novel human secreted and transmembrane protein PRO733.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
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06-NOV-2003.

(GETH ) GENENTECH INC.

76.0%; Score 174; DB 8; Lé

lery Match 76.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.

16-Y Match 76.0%; Score 174; DB 8; Langery Match 76.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Fred. No. 1.2e-158;
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PA (GETH ) GENENTECH INC.
QUERY MARCh 76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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US2003207353-A1.
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US2003207375-A1.
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Human PRO polypeptide #205.
US2003207425-Al.
                                                                                                               ADG03490 standard, protein, 229 AA Human PRO polypeptide #205. US2003207351-A1. G6-NOV-2003. (GETH ) GENENTECH INC. 76.0%; Scoresty Match
                                                                                                                                                                                                                                                                                                                                                                                                                 ADG16796 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003207359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG13359 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003207357-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG15586 standard; protein; 229 AA Human PRO polypeptide #205.
US2003219885-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF96984 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #205.
US2003207371-A1.
06-NOV-2003.
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Best Local Similarity
RESULT 253
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RESULT 251
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Best Local Similarity
RESULT 252
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Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 277

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DB 8; Length 229;
Length 229;
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Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207362-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG70965 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207420-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG58047 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG53631 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207415-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207421-A1.
                                                                                                                                                                                                                                                                                                                                                                                                  ADG56943 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG55839 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG58599 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207368-A1.
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                                                                                                                                                                                                                                                                                                                                       Score 174; DB 8; Lo
Pred. No. 1.2e-158;
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76.0%; Score 174; DB 8; L 100.0%; Pred. No. 1.2e-158;
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06-NOV-2003.
(GETH ) GENENTECH INC.
76.0%; SCORE 174;
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100.0%;
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                                                                                   Human PRO polypeptide #205.
US2003207358-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
Best Local Similarity
RESULT 268
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RESULT 270
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                                                               ADG82256 standard;
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RESULT 273
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vuery Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 281
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ry Match 76.0%; Score 174; DB 8; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH12033 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207419-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207414-A1.
ADG50620 standard; protein; 229 AA.

Human secreted/transmembrane protein, PRO733.
US2003207803-A1.
US-NOV-2003.
(GETH) GENENTECH INC.
T6.0%; Score 174; DB 8; Lest Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003215905-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, PRO733. US2001216305-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG51868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003215908-A1.
                                                                                                                                                                                                                                                                                                                         ADH30666 standard; protein; 229 AA. Human PRO polypeptide #205. 
US2003077723-A1. 
24-APR-2003.
                                                                                                                                                                       ADGB1704 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003207805-A1.
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24-APR-2003.
(GETH ) GENENTECH INC.
76.0%; S/
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(GETH ) GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
76.0%;
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                                                                                                                       Best Local Similarity RESULT 278
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DB 8; Length 229;

Length 229;

Length 229;

DB 8;

Length 229;

Length 229;

RESULT

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ADH25669 standard; protein; 229 AA.
Human neurotrimin homologue related protein sequence SEQ ID NO:447.
                                                                                                                                                                                                                                             76.0%; Score 174; DB 8; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADII15413 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207382-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human secreted and transmembrane protein PRO733. US2004009548-Al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADII4745 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207383-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI18340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207349-A1.
06-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ63621 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
                                                                                                                                                                                                                                                                                                                                                             PA (GETH) GENENTECH INC.
Query Match
76.0%; Score 174; DB 8; L4
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-JAN-2004.
(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 8; Le
ery Match 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ) 15-JAN-2004.

A (GETH) GENENTECH INC.

Query Match.

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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ry Match 76.0%; Score 174; DB 8; Lu
t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                     Score 174; DB 8; I
Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
Human secreted/transmembrane protein, PRO733.
US2004005657-A1.
08-JAN-2004.
                                                                                                                                            ADG62644 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004006219-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 174;
                                                                                                                                                                                                                                                                                                  ADI81199 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003207361-A1.
06-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 229 AA
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                          400565.
AN -2004.
H ) GENENTECH INC.
76.0%; SCC
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                                                                                                                                                                                                    08-JAN-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                               Local Similarity
                                                                                                     Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG09942 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG09290 standard;
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                                                                                       Query Match
                                                                                                                                                                                                                                             Query Match
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                                                                  (GETH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 301
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                                                      16-OCT-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207369-A1.
                                                                                                                                                                              Novel human secreted and transmembrane protein PRO733 US2003207366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG61503 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207429-A1.
                                                                                                                                                                                                                                                                                                                                          Novel human secreted and transmembrane protein PRO733. US2003207378-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG54735 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207367-A1.
                                                                                                                                                                                                                                                                                                                                                                     OG-NOV-2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; Le
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; L/
                                                                                                                                                                                                                                                               Score 174; DB 8; L. Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG48748 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003216560-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADGS1244 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004005312-A1.
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JOJAN-2003.

(GETH) GENENTECH INC.

(GETH) GENENTECH 176; Score 174;

PER SCORE 174;

ACT STATE SCORE 174;
    ADG81152 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194793-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO polypeptide #205.
                                                                                                                                                                protein; 229 AA.
                                                                                                                                                                                                                                                                                                                      ADH12657 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG59188 standard; protein; 229 AA
                                                                                                                                                                                                US200320...
06-NOV-2003.
(GETH ) GENENTECH INC.
76.0%; S
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2003.

2003.

241.

267 GENENTECH IN

LATY MATCH

Best Local Similarity 1v

RESULT 291

ID ADH28590 standar

DE Human PRO reference PN US2003r

PD 30-
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(GETH ) GENENTECH INC.
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2003.

2003.

21 Match
Best Local Similarity
RESULT 290
D ADG61503 stand*
DE Novel humar
PN US2003***
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                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 288
                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 289
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                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 295
                                                                                                                                                            ADG56391 standard;
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BEREE

Length 229;

DB 8; Length 229;

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(GETH) GENENTECH INC.
Query Match
76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                              14-OCT-2004.

(GETH) GENENTECH INC.

QUETY MATCh

76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 76.0%; Score 174; DB 8; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                               Novel bronchial cancer-associated human protein SeqID899.
DE10316701-A1.
                                                                                         ADS32346 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2004203125-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-NOV-2004.
(GETH) GENENTECH INC.
176.0%; Score 174; DB 8; Le
tt Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                / Match 76.0%; Score 174; DB 8; L. Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                           Human PRO polypeptide #205.
US2004214269-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADUS0402 standard; protein; 229 AA.
PRO733, SEQ ID 447.
US2004223964-A1.
                                                                                                                                                                                                                                                                                                                                                                                      protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HINZ/) HINZMANN B.
(HERM/) HERMANN K.
(CAST/) HEIDEN CASTANOS-VELEZ E.
                                                                                                                                                                                                                                                                                     CGETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRO733 protein, SEQ ID 447
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TUMAS D.
WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EATON D L.
FERRARA N.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ROY M A.
SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
RESULT 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DESNOYERS L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PAN J.
PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KUO S S.
NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                      ADU06673 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADW49681 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GODDARD A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GURNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GODOWSKI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GERBER
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     06-NOV-2003
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                                                                            RESULT 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 229;
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                                                                                                                                                                                                        Length 229;
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                                                        76.0%; Score 174; DB 8; Length 229; 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-Apr-2004.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; LE.
100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 25-MAR-2004.
PA (EETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                USJULAR-2004.

(GETH ) GENENTECH INC.

ALY MALCh ... 13 TILY 100.0%; Score 174; DB 8; L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 76.0%; Score 174; DB 8; L Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1.2e-158;
                                                                                                                                                                                       Query Match 76.0%; Score 174; DB 8; L Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 11-MAR-2004.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 309
                                                                                                                                                                                                                                                                                                                                       Query Match 76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL07280 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004063921-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein, PRO733 US2004048332-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76.0%; Score 174;
                                                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #205.
US2004048333-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #205.
24-app.
                                                                                                   Human PRO polypeptide #205.
US2004038336-Al.
                                                                                                                                                                                                                                                         ADJ65638 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004038335-A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM42498 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004058424-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM28360 standard; protein; 229 AA. Human PRO polypeptide #205.
US2004077064-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM17446 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.08;
US2004039164-A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                  26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-APR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 313
                                                                     Best Local Similarity RESULT 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 312
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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13-OCT-2005.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                            SMITH V.
STEWART T A.
TUMAS D.
WATANABE C K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GERRITSEN M E.
GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TUMAS D.
WATANABE C K.
WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DEFORGE L.
DESNOYERS L.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GODOWSKI P J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                 GURNEY A L. SHERWOOD S.
                                                                                                                                              Query Match
Best Local Similarity
RESULT 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHERWOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GURNEY A L.
       GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BAKE/) BAKER K P.
(BERE/) BERESINI M
                                                                                                                                  ZHANG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SMITH V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZHANG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-FEB-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A (WOOD/) WC
A (ZHAN/) ZH
Query Match
                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                     (MOOD/)
(ZHAN/)
                                                             (SMIT/)
(STEW/)
                                                                                         (TUMA/)
(WATA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TUMA/)
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(STEW/)
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(DESN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (FILV/)
(GAOW/)
                                                (SHER/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GURN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
RESULT 329
                                                                                                                                                                                                                                                                                                                                                                                            Best I
RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEB14127 standard; protein; 229 AA.
Cancer cell diagnosis method-related human protein - SEQ ID 410.
US2005153396-A1.
                                                                         Length 229;
                                                                                                                                                                                         Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76.0%; Score 174; DB 9; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 229;
                                 PD 24-FEB-2005.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 320
                                                                                                                                                           PD 07-APR-2005.

PA (GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 9; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     76.0%; Score 174; DB 9; L 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                               ADZ52342 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2005084935-A1.
                                                                                                               ADZ03381 standard; protein; 229 AA.
Human secreted/transmembrane PRO733 protein.
US2005074837-A1.
ADY14737 standard; protein; 229 AA.
PRO polypeptide SEQ ID NO 543.
WO2005016962-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AEA99614 standard, protein, 229 AA.
Human PRO733 protein SEQ ID NO:2.
US2005118635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JUN-2005.
(GODD/) GODDARD A.
(GODC/) GODOWSKI P J.
(GURNA/) SMITH V.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JO/)
JEMY/) GD...
SMITH
(WOOD/) WOOD W I...
LETY MATCH
JEMIL 323
LD AEB14127 Standard;
DE CANCEr Cell diagr
PN US2005153396-A)
PD 14-JUL-2005.
PA (BAKE/) BF
PA (BREK/) BF
PA (DEFO/)
PA (DEFO/)
PA (FII'.
PA (G')
                                                                                                                                                                                                                                                                                        ASHKENAZI A J.
BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TUMAS D.
WILLIAMS P M.
WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                     GERRITSEN M E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ROY M A. SHELTON D L. STEWART T A.
                                                                                                                                                                                                                                                                                                                   BOTSTEIN D.
DESNOYERS L.
EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                              FILVAROFF E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PAN J.
PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GURNEY A L.
                                                                                                                                                                                                                                                                                                                                                             PERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GODDARD A
                                                                                                                                                                                                                                                                                                                                                                                                                     GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KUO S S.
                                                                                                                                                                                                                                                                                                                                                                                                          GAO W.
                                                                                                                                                                                                                                                                                                      (BAKE/)
                                                                                                                                                                                                                                                                                                                    (BOTS/)
                                                                                                                                                                                                                                                                                                                                                               FERR/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAON/
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Extended human secreted protein sequence, SEQ ID NO. 149.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                        Ouery Match
                                                                                                                                                                                                                    Query Match
                                                                                           ABI17411 standard; protein; 229 AA.
Human cancer treatment biomarker amino acid sequence - SEQ ID 298.
US2006121511-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADL22684 standard; protein; 141 AA.
Human disease detection and treatment (MDDT) protein - SEQ ID 133.
WO2003062379-A2.
                                                                                                                                                                                                                                                                                                                                           21-JUN-2006.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 10; Length 229;
                                                   76.0%; Score 174; DB 10; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                         Length 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               56.3%; Score 129; DB 7; Length 229; 100.0%; Pred. No. 2.3e-115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 8; Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  68.6%; Score 157; DB 7; Length 162; 100.0%; Pred. No. 2e-142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 133; DB 7; Length 141; Pred. No. 2.2e-119;
                                                                                                                                                                                                                                                                                 69.9%; Score 160; DB 2; Length 229; 100.0%; Pred. No. 3.6e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 193;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             211
                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY35962 standard; protein; 229 AA.
Extended human secreted protein sequence, SEQ ID NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 3.6e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 157; DB 7; L. Pred. No. 2.4e-142;
                                                                                                                                                                                                                                     76.0%; Score 174; DB 10; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    69.9%; Score 160; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE95518 standard; protein; 162 AA.
Human NOVX4b protein.
WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG14988 standard; protein; 193 AA.
Human SECP-21 protein.
WO2003087300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE95516 standard, protein, 229 AA.
Human NOVX4a protein.
WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP19270 standard; protein; 229 AA. Human secreted polypeptide #121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY35900 standard; protein; 162 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INCY-) INCYTE GENOMICS INC. ry Match 58.1%; & t Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 338
                                                                                                                                                             08-JUN-2006.
(LEEH/) LEE H.
(SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 334
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RESULT 335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 333
                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 331
                                                                    Best Local Similarity
           (SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUN-1999.
(GEST ) GENSET.
(LEEH/) LEE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9931236-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUN-200
                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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AB023233 standard, protein; 169 AA.
Human breast tumour associated protein 47-like polypeptide NOV1
US2003027158-A1.
                                                                54.6%; Score 125; DB 2; Length 162. 100.0%; Pred. No. 1.2e-111;
                                                                                                                                                                                                                                 Length 162;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             34.9%; Score 80; DB 8; Length 169; 100.0%; Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 8; Length 169;
2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6; Length 169;
2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34.9%; Score 80; DB 7; Length 169; 100.0%; Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                 41.9%; Score 96; DB 2; Length 120; 100.0%; Pred. No. 7.2e-84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4; Length 169;
                                                                                                                                                                                                                                                                                         AAY35901 standard; protein; 120 AA.
Extended human secreted protein sequence, SEQ ID NO. 150.
WO9931236-A2.
                                                                                                                                                                                                                               54.6%; Score 125; DB 8; I 100.0%; Pred. No. 1.2e-111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41.9%; Score 96; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 34.9%; Score 80;
Best Local Similarity 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 80;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 34.9%; Score 80; Best Local Similarity 100.0%; Pred. No.
                                                                                                         Human secreted polypeptide #59. US2004110939-A1. 10-UN-2004. GEST ) GENSET SA. ry Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB61129 standard; protein; 169 AA.
Human NOV1 protein.
WO200075321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI19766 standard; protein; 169 AA.
Human NOV1 protein.
US2004002134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP19209 standard; protein; 120 AA.
Human secreted polypeptide #60.
US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM56365 standard; protein; 169 AA.
Human cell adhesion molecule NOVI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF66729 standard; protein; 169 AA.
Novel human protein NOV1.
US2003199103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADO60239 standard; protein; 169 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-2000.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-FEB-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-) CURAGEN CORP.
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(GEST ) GENSET.
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